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CA 2403370 A1 2001/09/15

(21) 2 403 370

(12) DEMANDE DE BREVET CANADIEN
CANADIAN PATENT APPLICATION

(13) A1

(86) Date de dépôt PCT/PCT Filing Date: 2001/03/15
(87) Date publication PCT/PCT Publication Date: 2001/09/15
(85) Entrée phase nationale/National Entry: 2002/09/16
(86) N° demande PCT/PCT Application No.: US 2001/008678
(87) N° publication PCT/PCT Publication No.: 2001/068859
(30) Priorités/Priorities: 2000/03/16 (60/189,816) US;
2000/11/28 (09/724,460) US

(51) Cl.Int.⁷/Int.Cl.⁷ C12N 15/12, A61K 31/70, A01K 67/027,
A61K 38/17, C07K 16/18, C12N 15/11, C12N 5/10,
C12N 15/86, C12N 15/62, C07K 14/47, C12N 1/21,
G01N 33/68
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(54) Titre : MOLECULES DU TYPE RECEPTEUR DE L'INTERLEUKINE 17 ET LEUR UTILISATION
(54) Title: IL-17 RECEPTOR LIKE MOLECULES AND USES THEREOF

(57) Abrégé/Abstract:

Novel IL-17 receptor like polypeptides and nucleic acid molecules encoding the same. The invention also provides vectors, host cells, agonists and antagonists (including selective binding agents), and method for producing IL-17 receptor like polypeptides. Also provided for are methods for the treatment, diagnosis, amelioration, or prevention of diseases with IL-17 receptor like polypeptides.

(12) INTERNATIONAL APPLICATION PUBLISHED UNDER THE PATENT COOPERATION TREATY (PCT)

(19) World Intellectual Property Organization
International Bureau



(43) International Publication Date
20 September 2001 (20.09.2001)

PCT

(10) International Publication Number
WO 01/68859 A3

- (51) International Patent Classification⁷: C12N 15/12, C07K 14/47, C12N 1/21, 5/10, G01N 33/68, A61K 38/17, 31/70, C07K 16/18, C12N 15/62, 15/86, A01K 67/027, C12N 15/11
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- (81) Designated States (*national*): AE, AG, AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, BZ, CA, CH, CN, CO, CR, CU, CZ, DE, DK, DM, DZ, EE, ES, FI, GB, GD, GE, GH, GM, HR, HU, ID, IL, IN, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MA, MD, MG, MK, MN, MW, MX, MZ, NO, NZ, PL, PT, RO, RU, SD, SE, SG, SI, SK, SL, TJ, TM, TR, TT, TZ, UA, UG, US, UZ, VN, YU, ZA, ZW.
- (21) International Application Number: PCT/US01/08678
- (22) International Filing Date: 15 March 2001 (15.03.2001)
- (25) Filing Language: English
- (26) Publication Language: English
- (30) Priority Data:
60/189,816 16 March 2000 (16.03.2000) US
09/724,460 28 November 2000 (28.11.2000) US
- (84) Designated States (*regional*): ARIPO patent (GH, GM, KE, LS, MW, MZ, SD, SL, SZ, TZ, UG, ZW), Eurasian patent (AM, AZ, BY, KG, KZ, MD, RU, TJ, TM), European patent (AT, BE, CH, CY, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE, TR), OAPI patent (BF, BJ, CF, CG, CI, CM, GA, GN, GW, ML, MR, NE, SN, TD, TG).

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Published:

— with international search report

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(88) Date of publication of the international search report:
21 March 2002

For two-letter codes and other abbreviations, refer to the "Guidance Notes on Codes and Abbreviations" appearing at the beginning of each regular issue of the PCT Gazette.

WO 01/68859 A3

(54) Title: IL-17 RECEPTOR LIKE MOLECULES AND USES THEREOF

(57) Abstract: Novel IL-17 receptor like polypeptides and nucleic acid molecules encoding the same. The invention also provides vectors, host cells, agonists and antagonists (including selective binding agents), and method for producing IL-17 receptor like polypeptides. Also provided for are methods for the treatment, diagnosis, amelioration, or prevention of diseases with IL-17 receptor like polypeptides.

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WHAT IS CLAIMED

1. An isolated nucleic acid molecule comprising a nucleotide sequence selected from the group consisting of:

5 (a) the nucleotide sequence as set forth in SEQ ID NO: 1;

(b) a nucleotide sequence encoding the polypeptide as set forth in SEQ ID NO: 2;

(c) a nucleotide sequence which hybridizes under moderately or highly stringent conditions to the complement of

10 (a) or (b), wherein the encoded polypeptide has an activity of the polypeptide as set forth in SEQ ID NO: 2; and

(d) a nucleotide sequence complementary to any of (a)-(c).

15 2. An isolated nucleic acid molecule comprising a nucleotide sequence selected from the group consisting of:

(a) a nucleotide sequence encoding a polypeptide that is at least about 70, 75, 80, 85, 90, 95, 96, 97, 98, or 99 percent identical to the polypeptide as set forth in SEQ ID NO: 2, wherein the polypeptide has an activity of the polypeptide as set forth in SEQ ID NO: 2;

20 (b) a nucleotide sequence encoding an allelic variant or splice variant of the nucleotide sequence as set forth in SEQ ID NO: 1, wherein the encoded polypeptide has an activity of the polypeptide as set forth in SEQ ID NO: 2;

(c) a nucleotide sequence of SEQ ID NO: 1; (a); or (b) encoding a polypeptide fragment of at least about 25 amino acid residues, wherein the polypeptide has an activity of the polypeptide as set forth in SEQ ID NO: 2;

25 (d) a nucleotide sequence of SEQ ID NO: 1, or (a)-(c) comprising a fragment of at least about 16 nucleotides;

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(e) a nucleotide sequence which hybridizes under moderately or highly stringent conditions to the complement of any of (a)-(d), wherein the polypeptide has an activity of the polypeptide as set forth in SEQ ID NO: 2; and

5 (f) a nucleotide sequence complementary to any of (a)-(c).

3. An isolated nucleic acid molecule comprising a nucleotide sequence selected from the group consisting of:

10 (a) a nucleotide sequence encoding a polypeptide as set forth in SEQ ID NO: 2 with at least one conservative amino acid substitution, wherein the polypeptide has an activity of the polypeptide as set forth in SEQ ID NO: 2;

15 (b) a nucleotide sequence encoding a polypeptide as set forth in SEQ ID NO: 2 with at least one amino acid insertion, wherein the polypeptide has an activity of the polypeptide as set forth in SEQ ID NO: 2;

20 (c) a nucleotide sequence encoding a polypeptide as set forth in SEQ ID NO: 2 with at least one amino acid deletion, wherein the polypeptide has an activity of the polypeptide as set forth in SEQ ID NO: 2;

25 (d) a nucleotide sequence encoding a polypeptide as set forth in SEQ ID NO: 2 which has a C- and/or N- terminal truncation, wherein the polypeptide has an activity of the polypeptide as set forth in SEQ ID NO: 2;

30 (e) a nucleotide sequence encoding a polypeptide as set forth in SEQ ID NO: 2 with at least one modification selected from the group consisting of amino acid substitutions, amino acid insertions, amino acid deletions, C-terminal truncation, and N-terminal truncation, wherein the polypeptide has an activity of the polypeptide as set forth in SEQ ID NO: 2;

(f) a nucleotide sequence of (a)-(e) comprising a fragment of at least about 16 nucleotides;

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(g) a nucleotide sequence which hybridizes under moderately or highly stringent conditions to the complement of any of (a)-(f), wherein the polypeptide has an activity of the polypeptide as set forth in SEQ ID NO: 2; and

5 (h) a nucleotide sequence complementary to any of (a)-(e).

4. A vector comprising the nucleic acid molecule of Claims 1, 2, or 3.

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5. A host cell comprising the vector of Claim 4.

6. The host cell of Claim 5 that is a eukaryotic cell.

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7. The host cell of Claim 5 that is a prokaryotic cell.

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8. A process of producing an IL-17 receptor like polypeptide comprising culturing the host cell of Claim 5 under suitable conditions to express the polypeptide, and optionally isolating the polypeptide from the culture.

9. A polypeptide produced by the process of Claim 8.

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10. The process of Claim 8, wherein the nucleic acid molecule comprises promoter DNA other than the promoter DNA for the native IL-17 receptor like polypeptide operatively linked to the DNA encoding the IL-17 receptor like polypeptide.

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11. The isolated nucleic acid molecule according to Claim 2 wherein the percent identity is determined using a computer program selected from the group consisting of GAP, BLASTP, BLASTN, FASTA, BLASTA, BLASTX, BestFit, and the Smith-Waterman algorithm.

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12. A process for determining whether a compound inhibits IL-17 receptor like polypeptide activity or production comprising exposing a cell according to Claims 5, 6, or 7 to the compound, and measuring IL-17 receptor like polypeptide 5 activity or production in said cell.

13. An isolated polypeptide comprising the amino acid sequence set forth in SEQ ID NO: 2.

10 14. An isolated polypeptide comprising the amino acid sequence selected from the group consisting of:

(a) an amino acid sequence for an ortholog of SEQ ID NO: 2, wherein the encoded polypeptide has an activity of the polypeptide as set forth in SEQ ID NO: 2;

15 (b) an amino acid sequence that is at least about 70, 80, 85, 90, 95, 96, 97, 98, or 99 percent identical to the amino acid sequence of SEQ ID NO: 2, wherein the polypeptide has an activity of the polypeptide as set forth in SEQ ID NO: 2;

20 (c) a fragment of the amino acid sequence set forth in SEQ ID NO: 2 comprising at least about 25 amino acid residues, wherein the polypeptide has an activity of the polypeptide as set forth in SEQ ID NO: 2;

25 (d) an amino acid sequence for an allelic variant or splice variant of either the amino acid sequence as set forth in SEQ ID NO: 2, or at least one of (a)-(b) wherein the polypeptide has an activity of the polypeptide as set forth in SEQ ID NO: 2.

15. An isolated polypeptide comprising the amino acid 30 sequence selected from the group consisting of:

(a) the amino acid sequence as set forth in SEQ ID NO: 2 with at least one conservative amino acid substitution, wherein the polypeptide has an activity of the polypeptide as set forth in SEQ ID NO: 2 as determined using the computer

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program of GAP, BLASTP, BLASTN, FASTA, BLASTA, BLASTX, BestFit, or the Smith-Waterman algorithm,

(b) the amino acid sequence as set forth in SEQ ID NO: 2 with at least one amino acid insertion, wherein the 5 polypeptide has an activity of the polypeptide as set forth in SEQ ID NO: 2;

(c) the amino acid sequence as set forth in SEQ ID NO: 2 with at least one amino acid deletion, wherein the polypeptide has an activity of the polypeptide as set forth in SEQ ID NO: 10 2;

(d) the amino acid sequence as set forth in SEQ ID NO: 2 which has a C- and/or N-terminal truncation, wherein the polypeptide has an activity of the polypeptide as set forth in SEQ ID NO: 2; and

15 (e) the amino acid sequence as set forth in SEQ ID NO: 2, with at least one modification selected from the group consisting of amino acid substitutions, amino acid insertions, amino acid deletions, C-terminal truncation, and N-terminal truncation, wherein the polypeptide has an activity of the 20 polypeptide as set forth in SEQ ID NO: 2.

16. An isolated polypeptide encoded by the nucleic acid molecule of Claims 1, 2, or 3.

25 17. The isolated polypeptide according to Claim 14 wherein the percent identity is determined using a computer program selected from the group consisting of GAP, BLASTP, BLASTN, FASTA, BLASTA, BLASTX, BestFit, and the Smith-Waterman algorithm.

30

18. A polypeptide according to claim 15 or 16 wherein the amino acid at position 45 of SEQ ID NO: 2 is glycine, proline, or alanine.

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19. A polypeptide according to claim 15 or 16 wherein the amino acid at position 227 of SEQ ID NO: 2 is phenylalanine, leucine, valine, isoleucine, alanine, or tyrosine.

5

20. A polypeptide according to claim 15 or 16 wherein the amino acid at position 363 of SEQ ID NO: 2 is serine, threonine, alanine or cysteine.

10 21. A polypeptide according to claim 15 or 16 wherein the amino acid at position 374 of SEQ ID NO: 2 is valine, isoleucine, methionine, leucine, phenylalanine, alanine or norleucine.

15 22. A polypeptide according to claim 15 or 16 wherein the amino acid at position 385 of SEQ ID NO: 2 is cysteine, serine, alanine.

20 23. A polypeptide according to claim 15 or 16 wherein the amino acid at position 515 of SEQ ID NO: 2 is aspartic acid or glutamic acid.

25 24. A polypeptide according to claim 15 or 16 wherein the amino acid at position 602 of SEQ ID NO: 2 is cysteine, alanine or serine.

25. An antibody produced by immunizing an animal with a peptide comprising an amino acid sequence of SEQ ID NO: 2.

30 26. An antibody or fragment thereof that specifically binds the polypeptide of Claims 13, 14, or 15.

27. The antibody of Claim 26 that is a monoclonal antibody.

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28. A hybridoma that produces a monoclonal antibody that binds to a peptide comprising an amino acid sequence of SEQ ID NO: 2.

5 29. A method of detecting or quantitating the amount of IL-17 receptor like polypeptide in a sample comprising contacting a sample suspected of containing IL-17 receptor like polypeptide with the anti-IL-17 receptor like antibody or fragment of Claims 25, 26, or 27 and detecting binding of said
10 antibody or fragment.

15 30. A selective binding agent or fragment thereof that specifically binds at least one polypeptide wherein said polypeptide comprises the amino acid sequence selected from the group consisting of:

- a) the amino acid sequence as set forth in SEQ ID NO: 2; and
- b) a fragment of the amino acid sequence set forth in at least one of SEQ ID NO: 2; and
- c) a naturally occurring variant thereof.

20 31. The selective binding agent of Claim 30 that is an antibody or fragment thereof.

25 32. The selective binding agent of Claim 30 that is a humanized antibody.

33. The selective binding agent of Claim 30 that is a human antibody or fragment thereof.

30 34. The selective binding agent of Claim 30 that is a polyclonal antibody or fragment thereof.

35. The selective binding agent of Claim 30 that is a monoclonal antibody or fragment thereof.

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36. The selective binding agent of Claim 30 that is a chimeric antibody or fragment thereof.

37. The selective binding agent of Claim 30 that is a
5 CDR-grafted antibody or fragment thereof.

38. The selective binding agent of Claim 30 that is an antiidiotypic antibody or fragment thereof.

10 39. The selective binding agent of Claim 30 which is a variable region fragment.

40. The variable region fragment of Claim 39 which is a Fab or a Fab' fragment.

15 41. A selective binding agent or fragment thereof comprising at least one complementarity determining region with specificity for a polypeptide having the amino acid sequence of SEQ ID NO: 2.

20 42. The selective binding agent of Claim 30 which is bound to a detectable label.

25 43. The selective binding agent of Claim 30 which antagonizes IL-17 receptor like polypeptide biological activity.

30 44. A method for treating, preventing, or ameliorating a disease, condition, or disorder associated with altered levels of IL-17 receptor like polypeptide comprising administering to a patient an effective amount of a selective binding agent according to Claim 30.

35 45. A selective binding agent produced by immunizing an animal with a polypeptide comprising an amino acid sequence

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selected from the group consisting of SEQ ID NO: 2.

46. A hybridoma that produces a selective binding agent capable of binding a polypeptide encoded by the nucleic acid
5 of Claims 1, 2, or 3.

47. A composition comprising the polypeptide of Claims 13, 14, or 15 and a pharmaceutically acceptable formulation agent.

10

48. The composition of Claim 47 wherein the pharmaceutically acceptable formulation agent is a carrier, adjuvant, solubilizer, stabilizer, or anti-oxidant or combination thereof.

15

49. The composition of Claim 47 wherein the polypeptide comprises the mature amino acid sequence as set forth in SEQ ID NO: 2.

20

50. A polypeptide comprising a derivative of the polypeptide of Claims 13, 14, or 15.

51. The polypeptide of Claim 50 which is covalently modified with a water-soluble polymer.

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52. The polypeptide of Claim 51 wherein the water-soluble polymer is selected from the group consisting of polyethylene glycol, monomethoxy-polyethylene glycol, dextran, cellulose, poly-(N-vinyl pyrrolidone) polyethylene glycol, propylene glycol homopolymers, polypropylene oxide/ethylene oxide co-polymers, polyoxyethylated polyols, and polyvinyl alcohol.
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53. A composition comprising a nucleic acid molecule of Claims 1, 2, or 3 and a pharmaceutically acceptable formulation agent.

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54. A composition of Claim 53 wherein said nucleic acid molecule is contained in a viral vector.

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55. A viral vector comprising a nucleic acid molecule of Claims 1, 2, or 3.

56. A fusion polypeptide comprising the polypeptide of
10 Claims 13, 14, or 15 fused to a heterologous amino acid sequence.

57. The fusion polypeptide of Claim 56 wherein the heterologous amino acid sequence is an IgG constant domain or
15 fragment thereof.

58. A method for treating, preventing or ameliorating a medical condition comprising administering to a patient the polypeptide of Claims 13, 14, or 15 or the polypeptide encoded
20 by the nucleic acid of Claims 1, 2, or 3 to said mammal.

59. A method of diagnosing a pathological condition or a susceptibility to a pathological condition in a subject caused by or resulting from abnormal levels of IL-17 receptor like
25 polypeptide comprising:

(a) determining the presence or amount of expression of the polypeptide of Claims 13, 14, or 15 or the polypeptide encoded by the nucleic acid molecule of Claims 1, 2, or 3 in a sample; and

30 (b) diagnosing a pathological condition or a susceptibility to a pathological condition based on the presence or amount of expression of the polypeptide.

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60. A device, comprising:

- (a) a membrane suitable for implantation; and
- (b) cells encapsulated within said membrane, wherein said cells secrete a protein of Claims 13, 14, or 15, and wherein
5 said membrane is permeable to said protein and impermeable to materials detrimental to said cells.

61. A device, comprising:

- (a) a membrane suitable for implantation; and
- 10 (b) the IL-17 receptor like polypeptide encapsulated within said membrane, wherein said membrane is permeable to the polypeptide.

62. A method of identifying a compound which binds to a
15 polypeptide comprising:

- (a) contacting the polypeptide of Claims 13, 14, or 15 with a compound; and
- (b) determining the extent of binding of the polypeptide to the compound.

20

63. A method of modulating levels of a polypeptide in an animal comprising administering to the animal the nucleic acid molecule of Claims 1, 2, or 3.

25 64. A transgenic non-human mammal comprising the nucleic acid molecule of Claims 1, 2, or 3.

30 65. A transgenic non-human comprising a disruption of the nucleic acid molecule of claim 1, 2 or 3 wherein the expression of IL-17 receptor polypeptide is decreased.

66. A method of identifying antagonists of IL-17 receptor like polypeptide biological activity comprising:

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- (a) contacting a compound with an IL-17 receptor like polypeptide;
- (b) detecting the biological activity of an IL-17 receptor like polypeptide in the presence of said compound;
- 5 and
- (c) comparing the level of IL-17 receptor like polypeptide biological activity in the presence and absence of said compound.

10 67. Then method of claim 66 wherein the compound is a small molecule, peptide, protein, carbohydrate, or antibody.

15 68. A method of modulating levels of a polypeptide in an animal comprising administering to the animal the nucleic acid molecule of claims 1, 2, or 3.

20 69. An antagonist of IL-17 receptor like polypeptide activity selected from the group consisting of IL-17 receptor like selective binding agents, small molecules, antisense oligonucleotides, and peptides or derivatives thereof having specificity for IL-17 receptor like polypeptide.

25 70. A method of reducing cellular production of IL-17 receptor like polypeptide, comprising transforming or transfected cells with a nucleic acid encoding an antagonist according to claim 69.

30 71. A method according to claim 70, wherein the antagonist is an antisense reagent, said reagent comprising an oligonucleotide comprising a single stranded nucleic acid sequence capable of binding to IL-17 receptor like mRNA.

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IL-17 RECEPTOR LIKE MOLECULES AND USES THEREOF

Related Application

5 The present application claims priority from U.S. patent application 09/724,460 file November 28, 2000 which claims priority from provisional application US 60/189,816 filed March 16, 2000.

10 Field of the Invention

The present invention relates to novel IL-17 receptor like polypeptides and nucleic acid molecules encoding the same. The invention also relates to vectors, host cells, pharmaceutical compositions, selective binding agents and methods for producing IL-17 receptor like polypeptides. Also provided for are methods for the diagnosis, treatment, amelioration, and/or prevention of diseases associated with IL-17 receptor like polypeptides.

20 Background of the Invention

Technical advances in the identification, cloning, expression and manipulation of nucleic acids have greatly accelerated the discovery of novel therapeutics based upon deciphering the human genome. Rapid nucleic acid sequencing techniques can now generate sequence information at unprecedented rates and, coupled with computational analyses, allow the assembly of overlapping sequences into partial and entire genomes and the identification of polypeptide-encoding regions. A comparison of a predicted amino acid sequence against a database compilation of known amino acid sequences can allow one to determine the extent of homology to previously identified sequences and/or structural landmarks. The cloning and expression of a polypeptide-encoding region of

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a nucleic acid molecule provides a polypeptide product for structural and functional analyses. The manipulation of nucleic acid molecules and encoded polypeptides to create variants and derivatives thereof may confer advantageous 5 properties on a product for use as a therapeutic.

In spite of the significant technical advances in genome research over the past decade, the potential for the development of novel therapeutics based on the human genome is still largely unrealized. Many genes encoding potentially 10 beneficial polypeptide therapeutics, or those encoding polypeptides, which may act as "targets" for therapeutic molecules, have still not been identified. In addition, structural and functional analyses of polypeptide products from many human genes have not been undertaken.

15 Accordingly, it is an object of the invention to identify novel polypeptides and nucleic acid molecules encoding the same, which have diagnostic or therapeutic benefit.

Summary of the Invention

20 The present invention relates to novel IL-17 receptor like nucleic acid molecules and encoded polypeptides.

The invention provides for an isolated nucleic acid molecule comprising a nucleotide sequence selected from the group consisting of:

25 (a) the nucleotide sequence as set forth in SEQ ID NO: 1;

(b) a nucleotide sequence encoding the polypeptide as set forth in SEQ ID NO: 2;

30 (c) a nucleotide sequence which hybridizes under moderately or highly stringent conditions to the complement of (a) or (b), wherein the encoded polypeptide has an activity of the polypeptide as set forth in SEQ ID NO: 2; and

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- (d) a nucleotide sequence complementary to any of (a)-(c).

5 The invention also provides for an isolated nucleic acid molecule comprising a nucleotide sequence selected from the group consisting of:

- (a) a nucleotide sequence encoding a polypeptide that is at least about 70, 75, 80, 85, 90, 95, 96, 97, 98, or 99 percent identical to the polypeptide as set forth in SEQ ID NO: 2, wherein the polypeptide has an activity of the polypeptide as set forth in SEQ ID NO: 2;
- (b) a nucleotide sequence encoding an allelic variant or splice variant of the nucleotide sequence as set forth in SEQ ID NO: 1, wherein the encoded polypeptide has an activity of the polypeptide as set forth in SEQ ID NO: 2;
- (c) a nucleotide sequence of SEQ ID NO: 1, (a), or (b) encoding a polypeptide fragment of at least about 25 amino acid residues, wherein the polypeptide has an activity of the polypeptide as set forth in SEQ ID NO: 2;
- (d) a nucleotide sequence of SEQ ID NO: 1, or (a)-(c) comprising a fragment of at least about 16 nucleotides;
- (e) a nucleotide sequence which hybridizes under moderately or highly stringent conditions to the complement of any of (a)-(d), wherein the polypeptide has an activity of the polypeptide as set forth in SEQ ID NO: 2; and
- (f) a nucleotide sequence complementary to any of (a)-(e).

30 The invention further provides for an isolated nucleic acid molecule comprising a nucleotide sequence selected from the group consisting of:

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- (a) a nucleotide sequence encoding a polypeptide as set forth in SEQ ID NO: 2 with at least one conservative amino acid substitution, wherein the polypeptide has an activity of the polypeptide as set forth in SEQ ID NO: 2;
- 5 (b) a nucleotide sequence encoding a polypeptide as set forth in SEQ ID NO: 2 with at least one amino acid insertion, wherein the polypeptide has an activity of the polypeptide as set forth in SEQ ID NO: 2;
- 10 (c) a nucleotide sequence encoding a polypeptide as set forth in SEQ ID NO: 2 with at least one amino acid deletion, wherein the polypeptide has an activity of the polypeptide as set forth in SEQ ID NO: 2;
- 15 (d) a nucleotide sequence encoding a polypeptide as set forth in SEQ ID NO: 2 which has a C- and/or N- terminal truncation, wherein the polypeptide has an activity of the polypeptide as set forth in SEQ ID NO: 2;
- 20 (e) a nucleotide sequence encoding a polypeptide as set forth in SEQ ID NO: 2 with at least one modification selected from the group consisting of amino acid substitutions, amino acid insertions, amino acid deletions, C-terminal truncation, and N-terminal truncation, wherein the polypeptide has an activity of the polypeptide as set forth in SEQ ID NO: 2;
- (f) a nucleotide sequence of (a)-(e) comprising a fragment of at least about 16 nucleotides;
- 25 (g) a nucleotide sequence which hybridizes under moderately or highly stringent conditions to the complement of any of (a)-(f), wherein the polypeptide has an activity of the polypeptide as set forth in SEQ ID NO: 2; and
- (h) a nucleotide sequence complementary to any of (a)-
30 (e).

- 5 -

The invention also provides for an isolated polypeptide comprising the amino acid sequence selected from the group consisting of:

(a) an amino acid sequence for an ortholog of SEQ ID NO: 5 2, wherein the encoded polypeptide has an activity of the polypeptide as set forth in SEQ ID NO: 2;

(b) an amino acid sequence that is at least about 70, 80, 85, 90, 95, 96, 97, 98, or 99 percent identical to the amino acid sequence of SEQ ID NO: 2, wherein the polypeptide has an 10 activity of the polypeptide as set forth in SEQ ID NO: 2 as determined using a computer program such as GAP, BLASTP, BLASTN, FASTA, BLASTA, BLASTX Bestfit, and the Smith-Waterman algorithm;

(c) a fragment of the amino acid sequence set forth in 15 SEQ ID NO: 2 comprising at least about 25 amino acid residues, wherein the polypeptide has an activity of the polypeptide as set forth in SEQ ID NO: 2;

(d) an amino acid sequence for an allelic variant or splice variant of either the amino acid sequence as set forth 20 in SEQ ID NO: 2, or at least one of (a)-(b) wherein the polypeptide has an activity of the polypeptide as set forth in SEQ ID NO: 2.

The invention further provides for an isolated 25 polypeptide comprising the amino acid sequence selected from the group consisting of:

(a) the amino acid sequence as set forth in SEQ ID NO: 2 with at least one conservative amino acid substitution, wherein the polypeptide has an activity of the polypeptide as 30 set forth in SEQ ID NO: 2;

(b) the amino acid sequence as set forth in SEQ ID NO: 2 with at least one amino acid insertion, wherein the polypeptide has an activity of the polypeptide as set forth in SEQ ID NO: 2;

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(c) the amino acid sequence as set forth in SEQ ID NO: 2 with at least one amino acid deletion, wherein the polypeptide has an activity of the polypeptide as set forth in SEQ ID NO: 2;

5 (d) the amino acid sequence as set forth in SEQ ID NO: 2 which has a C- and/or N-terminal truncation, wherein the polypeptide has an activity of the polypeptide as set forth in SEQ ID NO: 2; and

10 (e) the amino acid sequence as set forth in SEQ ID NO: 2, with at least one modification selected from the group consisting of amino acid substitutions, amino acid insertions, amino acid deletions, C-terminal truncation, and N-terminal truncation, wherein the polypeptide has an activity of the polypeptide as set forth in SEQ ID NO: 2.

15 Also provided are fusion polypeptides comprising the amino acid sequences of (a)-(e) above.

The present invention also provides for an expression vector comprising the isolated nucleic acid molecules as set forth herein, recombinant host cells comprising recombinant nucleic acid molecules as set forth herein, and a method of producing an IL-17 receptor like polypeptide comprising culturing the host cells and optionally isolating the polypeptide so produced. These expression vectors include baculovirus expression vectors which utilize insect cells for expression.

A transgenic non-human animal comprising a nucleic acid molecule encoding an IL-17 receptor like polypeptide is also encompassed by the invention. The IL-17 receptor like nucleic acid molecules are introduced into the animal in a manner that allows expression and increased levels of the IL-17 receptor like polypeptide, which may include increased circulating levels. The transgenic non-human animal is preferably a mammal. Also provided is a transgenic non-human animals

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comprising a disruption in the nucleic acid molecule encoding a IL-17 receptor like polypeptide, which will knock-out or specifically decrease expression of the IL-17 receptor like polypeptide.

5 Also provided are derivatives of the IL-17 receptor like polypeptides of the present invention.

Analogs of the IL-17 receptor like polypeptides are provided for in the present invention which result from conservative and/or non-conservative amino acids substitutions 10 of the IL-17 receptor like polypeptides of SEQ ID NO: 2. Such analogs include an IL-17 receptor like polypeptide wherein, for example the amino acid at position 45 of SEQ ID NO: 2 is glycine, proline or alanine, the amino acid at position 227 of SEQ ID NO: 2 is phenylalanine, leucine, valine, isoleucine, 15 alanine or tyrosine, the amino acid at position 363 of SEQ ID NO: 2 is serine, threonine, alanine or cysteine, the amino acid at position 374 of SEQ ID NO: 2 is valine, isoleucine, leucine, phenylalanine, alanine or norleucine, the amino acid at position 515 of SEQ ID NO: 2 is aspartic acid or glutamic 20 acid, the amino acid at position 6002 of SEQ ID NO: 2 is cysteine, serine or alanine.

Additionally provided are selective binding agents such as antibodies and peptides capable of specifically binding the IL-17 receptor like polypeptides of the invention. Such 25 antibodies, polypeptides, peptides and small molecules may be agonistic or antagonistic.

Pharmaceutical compositions comprising the nucleotides, polypeptides, or selective binding agents of the present invention and one or more pharmaceutically acceptable 30 formulation agents are also encompassed by the invention. The pharmaceutical compositions are used to provide therapeutically effective amounts of the nucleotides or polypeptides of the present invention. The invention is also

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directed to methods of using the polypeptides, nucleic acid molecules, and selective binding agents.

The IL-17 receptor like polypeptides and nucleic acid molecules of the present invention may be used to treat, 5 prevent, ameliorate, diagnose and/or detect diseases and disorders, including those recited herein. Expression analysis in biological, cellular, or tissue samples suggests that IL-17 receptor like polypeptide may play a role in the diagnosis and/or treatment of the pathological conditions 10 described herein. This expression can be detected with a diagnostic agent such as a IL-17 receptor like polynucleotide.

The invention encompasses diagnosing a pathological condition or the susceptibility to a pathological condition in a subject caused by or resulting from abnormal (i.e. increased 15 or decreased) levels of IL-17 receptor like polypeptide comprising determining the presence or amount of expression of the IL-17 receptor like polypeptide in a sample and comparing the level of said polypeptide in a biological, tissue or cellular sample form either normal subjects at an earlier 20 time, wherein susceptibility to a a pathological condition is based on the presence or amount of expression of the polypeptide.

The present invention also provides a method of assaying test molecules to identify a test molecule which binds to an 25 IL-17 receptor like polypeptide. The method comprises contacting an IL-17 receptor like polypeptide with a test molecule and determining the extent of binding of the test molecule to the polypeptide. The method further comprises determining whether such test molecules are agonists or 30 antagonists of an IL-17 receptor like polypeptide. The present invention further provides a method of testing the impact of molecules on the expression of IL-17 receptor like polypeptide or on the activity of IL-17 receptor like polypeptide.

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The present invention provides for methods of identifying antagonists and agonists of IL-17 receptor like biological activity comprising contacting a small molecule compound with IL-17 receptor like polypeptide and measuring IL-17 receptor like biological activity in the presence and absence of these small molecules. These small molecules can be a naturally occurring medicinal compound derived from combinatorial chemical libraries. In certain embodiments, an IL-17 receptor like polypeptide agonist or antagonist may be a protein, peptide, carbohydrate, lipid, or small molecule which interacts with a IL-17 receptor like polypeptide to regulate its activity.

Methods of regulating expression and modulating (i.e., increasing or decreasing) levels of an IL-17 receptor like polypeptide are also encompassed by the invention. One method comprises administering to an animal a nucleic acid molecule encoding an IL-17 receptor like polypeptide. In another method, a nucleic acid molecule comprising elements that regulate or modulate the expression of an IL-17 receptor like polypeptide may be administered. Examples of these methods include gene therapy, cell therapy, and anti-sense therapy as further described herein.

The IL-17 receptor like polypeptide can be used for identifying ligands thereof. Various forms of "expression cloning" have been used for cloning ligands for receptors. See e.g., Davis et al., *Cell*, 87:1161-1169 (1996). These and other IL-17 receptor like ligand cloning experiments are described in greater detail herein. Isolation of the IL-17 receptor like ligand(s) allows for the identification or development of novel agonists and/or antagonists of the IL-17 receptor like signaling pathway.

Agonists and antagonists include, but are not limited to, ligands to the IL-17 receptor like polypeptides, soluble IL-17

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receptor like polypeptides, anti-IL-17 receptor like selective binding agents (such as antibodies and derivatives thereof), small molecules, peptides and derivatives thereof, and antisense oligonucleotides, any of which can be used for
5 treating one or more disease or disorder, including those disclosed herein.

The invention further encompasses methods for determining the presence of IL-17 receptor like nucleic acids in a biological, tissue or cellular sample. These methods comprise
10 the steps of providing a biological sample suspected of containing IL-17 receptor like nucleic acids; contacting the sample with a diagnostic reagent of the present invention; detecting hybridization between nucleic acid in the biological sample and the diagnostic reagent; and comparing the level of
15 hybridization between the biological sample with the level of hybridization between a known concentration of IL-17 receptor like nucleic acid and the diagnostic reagent. The polynucleotide detected in these methods may be an IL-17 receptor like DNA and/or a IL-17 receptor like RNA.

20 The invention also provides for a device which comprises a membrane for implantation in a patient; and cells encapsulated within said membrane, wherein said membrane is permeable to the protein product and impermeable to materials detrimental to said cells. The invention further provides for
25 a device which comprises a membrane suitable for implantation and the IL-17 receptor like polypeptide encapsulated in a membrane that is permeable to the polypeptide.

Brief Description of the Figures

Figure 1 depicts a nucleic acid sequence (SEQ ID NO:1)
30 and amino acid sequence (SEQ ID NO: 2) of the human IL-17 receptor like polypeptide.

Figure 2 depicts an overlap of the present human IL-17 receptor like (hIL-17RL) polypeptide and a known IL-17

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receptor family member (SEQ ID NO: 3).

Detailed Description of the Invention

The section headings used herein are for organizational
5 purposes only and are not to be construed as limiting the
subject matter described therein. All references cited in
this application are expressly incorporated by reference
herein.

10 Definitions

The terms "IL-17 receptor like gene" or "IL-17 receptor
like nucleic acid molecule" or "polynucleotide" refers to a
nucleic acid molecule comprising or consisting of a nucleotide
sequence as set forth in SEQ ID NO: 1, a nucleotide sequence
15 encoding the polypeptide as set forth in SEQ ID NO: 2, a
nucleotide sequence of the DNA insert in Amgen deposit no. A-
672-P, and nucleic acid molecules as defined herein.

The term "IL-17 receptor like polypeptide" refers to a
polypeptide comprising the amino acid sequence of SEQ ID NO:
20 2, and related polypeptides. Related polypeptides include:
IL-17 receptor like polypeptide allelic variants, IL-17
receptor like polypeptide orthologs, IL-17 receptor like
polypeptide splice variants, IL-17 receptor like polypeptide
variants and IL-17 receptor like polypeptide derivatives. IL-
25 17 receptor like polypeptides may be mature polypeptides, as
defined herein, and may or may not have an amino terminal
methionine residue, depending on the method by which they are
prepared.

The term "IL-17 receptor like polypeptide allelic
30 variant" refers to one of several possible naturally occurring
alternate forms of a gene occupying a given locus on a
chromosome of an organism or a population of organisms.

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The term "IL-17 receptor like polypeptide derivatives" refers to the polypeptide as set forth in SEQ ID NO: 2, IL-17 receptor like polypeptide allelic variants, IL-17 receptor like polypeptide orthologs, IL-17 receptor like polypeptide splice variants, or IL-17 receptor like polypeptide variants, 5 as defined herein, that have been chemically modified.

The term "IL-17 receptor like polypeptide fragment" refers to a polypeptide that comprises a truncation at the amino terminus (with or without a leader sequence) and/or a 10 truncation at the carboxy terminus of the polypeptide as set forth in SEQ ID NO: 2, IL-17 receptor like polypeptide allelic variants, IL-17 receptor like polypeptide orthologs, IL-17 receptor like polypeptide splice variants and/or an IL-17 receptor like polypeptide variant having one or more amino 15 acid additions or substitutions or internal deletions (wherein the resulting polypeptide is at least 6 amino acids or more in length) as compared to the IL-17 receptor like polypeptide amino acid sequence set forth in SEQ ID NO: 2. IL-17 receptor like polypeptide fragments may result from alternative RNA 20 splicing or from *in vivo* protease activity. For transmembrane or membrane-bound forms of an IL-17 receptor like polypeptide, preferred fragments include soluble forms such as those lacking a transmembrane or membrane-binding domain. In preferred embodiments, truncations comprise about 10 amino 25 acids, or about 20 amino acids, or about 50 amino acids, or about 75 amino acids, or about 100 amino acids, or more than about 100 amino acids. The polypeptide fragments so produced will comprise about 25 contiguous amino acids, or about 50 amino acids, or about 75 amino acids, or about 100 amino acids, 30 or about 150 amino acids, or about 200 amino acids. Such IL-17 receptor like polypeptide fragments may optionally comprise an amino terminal methionine residue. It will be appreciated that such fragments can be used, for example, to generate antibodies to IL-17 receptor like polypeptides.

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The term "IL-17 receptor like fusion polypeptide" refers to a fusion of one or more amino acids (such as a heterologous peptide or polypeptide) at the amino or carboxy terminus of the polypeptide as set forth in SEQ ID NO: 2, IL-17 receptor like polypeptide allelic variants, IL-17 receptor like polypeptide splice variants, or IL-17 receptor like polypeptide variants having one or more amino acid deletions, substitutions or internal additions as compared to the IL-17 receptor like polypeptide amino acid sequence set forth in SEQ ID NO: 2.

The term "IL-17 receptor like polypeptide ortholog" refers to a polypeptide from another species that corresponds to IL-17 receptor like polypeptide amino acid sequence as set forth in SEQ ID NO: 2. For example, mouse and human IL-17 receptor like polypeptides are considered orthologs of each other.

The term "IL-17 receptor like polypeptide splice variant" refers to a nucleic acid molecule, usually RNA, which is generated by alternative processing of intron sequences in an RNA transcript of IL-17 receptor like polypeptide amino acid sequence as set forth in SEQ ID NO: 2.

The term "IL-17 receptor like polypeptide variants" refers to IL-17 receptor like polypeptides comprising amino acid sequences having one or more amino acid sequence substitutions, deletions (such as internal deletions and/or IL-17 receptor like polypeptide fragments), and/or additions (such as internal additions and/or IL-17 receptor like fusion polypeptides) as compared to the IL-17 receptor like polypeptide amino acid sequence set forth in SEQ ID NO: 2 (with or without a leader sequence). Variants may be naturally occurring (e.g., IL-17 receptor like polypeptide allelic variants, IL-17 receptor like polypeptide orthologs and IL-17 receptor like polypeptide splice variants) or

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artificially constructed. Such IL-17 receptor like polypeptide variants may be prepared from the corresponding nucleic acid molecules having a DNA sequence that varies accordingly from the DNA sequence as set forth in SEQ ID NO:

5 1. In preferred embodiments, the variants have from 1 to 3, or from 1 to 5, or from 1 to 10, or from 1 to 15, or from 1 to 20, or from 1 to 25, or from 1 to 50, or from 1 to 75, or from 1 to 100, or more than 100 amino acid substitutions, insertions, additions and/or deletions, wherein the

10 substitutions may be conservative, or non-conservative, or any combination thereof.

The term "antigen" refers to a molecule or a portion of a molecule capable of being bound by a selective binding agent, such as an antibody, and additionally capable of being used in an animal to produce antibodies capable of binding to an epitope of that antigen. An antigen may have one or more epitopes.

The term specific binding reaction referred to above is meant to indicate that the antigen will react, in a highly selective manner, with its corresponding antibody and not with the multitude of other antibodies which can be evoked by other antigens.

The term "biologically active IL-17 receptor like polypeptides" refers to IL-17 receptor like polypeptides, including fragments, variants, derivatives, having at least one activity characteristic of the polypeptide comprising the amino acid sequence of SEQ ID NO: 2. In addition, a IL-17 receptor like polypeptide may be active as an immunogen, that is, the polypeptide contains at least one epitope to which antibodies may be raised.

The terms "effective amount" and "therapeutically effective amount" each refer to the amount of a IL-17 receptor like polypeptide or IL-17 receptor like nucleic acid molecule

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used to support an observable level of one or more biological activities of the IL-17 receptor like polypeptides as set forth herein.

The term "expression vector" refers to a vector which is
5 suitable for use in a host cell and contains nucleic acid sequences which direct and/or control the expression of inserted heterologous nucleic acid sequences. Expression includes, but is not limited to, processes such as transcription, translation, and RNA splicing, if introns are
10 present.

The term "host cell" is used to refer to a cell which has been transformed, or is capable of being transformed with a nucleic acid sequence and then of expressing a selected gene of interest. The term includes the progeny of the parent
15 cell, whether or not the progeny is identical in morphology or in genetic make-up to the original parent, so long as the selected gene is present.

The term "identity" as known in the art, refers to a relationship between the sequences of two or more polypeptide molecules or two or more nucleic acid molecules, as determined by comparing the sequences. In the art, "identity" also means the degree of sequence relatedness between nucleic acid molecules or polypeptides, as the case may be, as determined by the match between strings of two or more nucleotide or two
20 or more amino acid sequences. "Identity" measures the percent of identical matches between the smaller of two or more sequences with gap alignments (if any) addressed by a particular mathematical model or computer program (i.e.,
25 "algorithms").

30 The term "similarity" is a related concept, but in contrast to "identity", refers to a measure of similarity which includes both identical matches and conservative substitution matches. If two polypeptide sequences have, for

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example, 10/20 identical amino acids, and the remainder are all non-conservative substitutions, then the percent identity and similarity would both be 50%. If in the same example, there are 5 more positions where there are conservative 5 substitutions, then the percent identity remains 50%, but the per cent similarity would be 75% (15/20). Therefore, in cases where there are conservative substitutions, the degree of similarity between two polypeptides will be higher than the percent identity between those two polypeptides.

10 The term "isolated nucleic acid molecule" refers to a nucleic acid molecule of the invention that (1) has been separated from at least about 50 percent of proteins, lipids, carbohydrates or other materials with which it is naturally found when total DNA is isolated from the source cells, (2) is 15 not linked to all or a portion of a polynucleotide to which the "isolated nucleic acid molecule" is linked in nature, (3) is operably linked to a polynucleotide which it is not linked to in nature, or (4) does not occur in nature as part of a larger polynucleotide sequence. Preferably, the isolated 20 nucleic acid molecule of the present invention is substantially free from at least one contaminating nucleic acid molecule with which it is naturally associated. Preferably, the isolated nucleic acid molecule of the present invention is substantially free from any other contaminating 25 nucleic acid molecule(s) or other contaminants that are found in its natural environment that would interfere with its use in polypeptide production or its therapeutic, diagnostic, prophylactic or research use.

30 The term "isolated polypeptide" refers to a polypeptide of the present invention that (1) has been separated from at least about 50 percent of polynucleotides, lipids, carbohydrates or other materials with which it is naturally found when isolated from the source cell, (2) is not linked (by covalent or noncovalent interaction) to all or a portion

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of a polypeptide to which the "isolated polypeptide" is linked to in nature, (3) is operably linked (by covalent or noncovalent interaction) to a polypeptide with which it is not linked in nature, or (4) does not occur in nature.

5 Preferably, is free from at least one contaminating polypeptide or other contaminants that are found in its natural environment. Preferably, the isolated polypeptide is substantially free from any other contaminating polypeptides or other contaminants that are found in its natural

10 environment which would interfere with its therapeutic, diagnostic, prophylactic or research use.

The term "mature IL-17 receptor like polypeptide" refers to an IL-17 receptor like polypeptide lacking a leader sequence. A mature IL-17 receptor like polypeptide may also

15 include other modifications such as proteolytic processing of the amino terminus (with or without a leader sequence) and/or the carboxy terminus, cleavage of a smaller polypeptide from a larger precursor, N-linked and/or O-linked glycosylation, and the like.

20 The term "nucleic acid sequence" or "nucleic acid molecule" refers to a DNA or RNA sequence. The term encompasses molecules formed from any of the known base analogs of DNA and RNA such as, but not limited to 4-acetylcytosine, 8-hydroxy-N6-methyladenosine, aziridinyl-

25 cytosine, pseudouracil, 5-(carboxyhydroxymethyl) uracil, 5-fluorouracil, 5-bromouracil, 5-carboxymethylaminomethyl-2-thiouracil, 5-carboxy-methylaminomethyluracil, dihydrouracil, inosine, N6-iso-pentenyladenine, 1-methyladenine, 1-methylpseudouracil, 1-methylguanine, 1-methylinosine, 2,2-dimethyl-guanine, 2-methyladenine, 2-methylguanine, 3-methylcytosine, 5-methylcytosine, N6-methyladenine, 7-methylguanine, 5-methylaminomethyluracil, 5-methoxyamino-methyl-2-thiouracil, beta-D-mannosylqueosine, 5'-methoxycarbonyl-methyluracil, 5-methoxyuracil, 2-methylthio-

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N6-isopentenyladenine, uracil-5-oxyacetic acid methylester,
uracil-5-oxyacetic acid, oxybutoxosine, pseudouracil,
queosine, 2-thiocytosine, 5-methyl-2-thiouracil, 2-thiouracil,
4-thiouracil, 5-methyluracil, N-uracil-5-oxyacetic acid
5 methylester, uracil-5-oxyacetic acid, pseudouracil, queosine,
2-thiocytosine, and 2,6-diaminopurine.

The term "naturally occurring" or "native" when used in connection with biological materials such as nucleic acid molecules, polypeptides, host cells, and the like, refers to materials which are found in nature and are not manipulated by man. Similarly, "non-naturally occurring" or "non-native" as used herein refers to a material that is not found in nature or that has been structurally modified or synthesized by man.

The term "operably linked" is used herein to refer to an arrangement of flanking sequences wherein the flanking sequences so described are configured or assembled so as to perform their usual function. Thus, a flanking sequence operably linked to a coding sequence may be capable of effecting the replication, transcription and/or translation of the coding sequence. For example, a coding sequence is operably linked to a promoter when the promoter is capable of directing transcription of that coding sequence. A flanking sequence need not be contiguous with the coding sequence, so long as it functions correctly. Thus, for example, intervening untranslated yet transcribed sequences can be present between a promoter sequence and the coding sequence and the promoter sequence can still be considered "operably linked" to the coding sequence.

The term "pharmaceutically acceptable carrier" or "physiologically acceptable carrier" as used herein refers to one or more formulation materials suitable for accomplishing or enhancing the delivery of the IL-17 receptor like polypeptide, IL-17 receptor like nucleic acid molecule or IL-

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17 receptor like selective binding agent as a pharmaceutical composition.

The term "selective binding agent" refers to a molecule or molecules having specificity for an IL-17 receptor like polypeptide. Selective binding agents include antibodies, such as polyclonal antibodies, monoclonal antibodies (mAbs), chimeric antibodies, CDR-grafted antibodies, anti-idiotypic (anti-Id) antibodies to antibodies that can be labeled in soluble or bound forms, as well as fragments, regions, or derivatives thereof which are provided by known techniques, including, but not limited to enzymatic cleavage, peptide synthesis, or recombinant techniques. The anti-IL-17 receptor like selective binding agents of the present invention are capable, for example, of binding portions of IL-17 like molecules to IL-17 receptor like polypeptides of the present invention.

As used herein, the terms, "specific" and "specificity" refer to the ability of the selective binding agents to bind to human IL-17 receptor like polypeptides and not to bind to human non-IL-17 receptor like polypeptides. It will be appreciated, however, that the selective binding agents may also bind orthologs of the polypeptide as set forth in SEQ ID NO: 2, that is, interspecies versions thereof, such as mouse and rat polypeptides.

25 The term "transduction" is used to refer to the transfer of genes from one bacterium to another, usually by a phage. "Transduction" also refers to the acquisition and transfer of eukaryotic cellular sequences by retroviruses.

30 The term "transfection" is used to refer to the uptake of foreign or exogenous DNA by a cell, and a cell has been "transfected" when the exogenous DNA has been introduced inside the cell membrane. A number of transfection techniques are well known in the art and are disclosed herein. See, for

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example, Graham et al., *Virology*, 52:456 (1973); Sambrook et al., *Molecular Cloning, a laboratory Manual*, Cold Spring Harbor Laboratories (New York, 1989); Davis et al., *Basic Methods in Molecular Biology*, Elsevier, 1986; and Chu et al., 5 *Gene*, 13:197 (1981). Such techniques can be used to introduce one or more exogenous DNA moieties into suitable host cells.

The term "transformation" as used herein refers to a change in a cell's genetic characteristics, and a cell has been transformed when it has been modified to contain a new 10 DNA. For example, a cell is transformed where it is genetically modified from its native state. Following transfection or transduction, the transforming DNA may recombine with that of the cell by physically integrating into a chromosome of the cell, may be maintained transiently as an 15 episomal element without being replicated, or may replicate independently as a plasmid. A cell is considered to have been stably transformed when the DNA is replicated with the division of the cell.

The term "vector" is used to refer to any molecule (e.g., 20 nucleic acid, plasmid, or virus) used to transfer coding information to a host cell.

Relatedness of Nucleic Acid Molecules
and/or Polypeptides

It is understood that related nucleic acid molecules 25 include allelic or splice variants of the nucleic acid molecule of SEQ ID NO: 1, and include sequences which are complementary to any of the above nucleotide sequences. Related nucleic acid molecules also include a nucleotide sequence encoding a polypeptide comprising or consisting 30 essentially of a substitution, modification, addition and/or a deletion of one or more amino acid residues compared to the polypeptide in SEQ ID NO: 2.

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Fragments include molecules which encode a polypeptide of at least about 25 amino acid residues, or about 50, or about 75, or about 100, or greater than about 100 amino acid residues of the polypeptide of SEQ ID NO: 2.

5 In addition, related IL-17 receptor like nucleic acid molecules include those molecules which comprise nucleotide sequences which hybridize under moderately or highly stringent conditions as defined herein with the fully complementary sequence of the nucleic acid molecule of SEQ ID NO: 1, or of a
10 molecule encoding a polypeptide, which polypeptide comprises the amino acid sequence as shown in SEQ ID NO: 2, or of a nucleic acid fragment as defined herein, or of a nucleic acid fragment encoding a polypeptide as defined herein.
Hybridization probes may be prepared using the IL-17 receptor
15 like sequences provided herein to screen cDNA, genomic or synthetic DNA libraries for related sequences. Regions of the DNA and/or amino acid sequence of IL-17 receptor like polypeptide that exhibit significant identity to known sequences are readily determined using sequence alignment
20 algorithms as described herein and those regions may be used to design probes for screening.

The term "highly stringent conditions" refers to those conditions that are designed to permit hybridization of DNA strands whose sequences are highly complementary, and to
25 exclude hybridization of significantly mismatched DNAs. Hybridization stringency is principally determined by temperature, ionic strength, and the concentration of denaturing agents such as formamide. Examples of "highly stringent conditions" for hybridization and washing are 0.015M sodium chloride, 0.0015M sodium citrate at 65-68°C or 0.015M sodium chloride, 0.0015M sodium citrate, and 50% formamide at 42°C. See Sambrook, Fritsch & Maniatis, Molecular Cloning: A Laboratory Manual, 2nd Ed., Cold Spring Harbor Laboratory, (Cold Spring Harbor, N.Y. 1989); Anderson *et al.*, Nucleic Acid

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Hybridisation: a practical approach, Ch. 4, IRL Press Limited (Oxford, England).

More stringent conditions (such as higher temperature, lower ionic strength, higher formamide, or other denaturing agent) may also be used, however, the rate of hybridization will be affected. Other agents may be included in the hybridization and washing buffers for the purpose of reducing non-specific and/or background hybridization. Examples are 0.1% bovine serum albumin, 0.1% polyvinyl-pyrrolidone, 0.1% sodium pyrophosphate, 0.1% sodium dodecylsulfate (NaDODSO_4 or SDS), ficoll, Denhardt's solution, sonicated salmon sperm DNA (or other non-complementary DNA), and dextran sulfate, although other suitable agents can also be used. The concentration and types of these additives can be changed without substantially affecting the stringency of the hybridization conditions. Hybridization experiments are usually carried out at pH 6.8-7.4, however, at typical ionic strength conditions, the rate of hybridization is nearly independent of pH. See Anderson et al., Nucleic Acid Hybridisation: a Practical Approach, Ch. 4, IRL Press Limited (Oxford, England).

Factors affecting the stability of a DNA duplex include base composition, length, and degree of base pair mismatch. Hybridization conditions can be adjusted by one skilled in the art in order to accommodate these variables and allow DNAs of different sequence relatedness to form hybrids. The melting temperature of a perfectly matched DNA duplex can be estimated by the following equation:

$$T_m(\text{°C}) = 81.5 + 16.6(\log[\text{Na}^+]) + 0.41(\%G+C) - 600/N - 0.72(\%\text{formamide})$$

where N is the length of the duplex formed, $[\text{Na}^+]$ is the molar concentration of the sodium ion in the hybridization or washing solution, %G+C is the percentage of (guanine+cytosine) bases in the hybrid. For imperfectly matched hybrids, the

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melting temperature is reduced by approximately 1°C for each 1% mismatch.

The term "moderately stringent conditions" refers to conditions under which a DNA duplex with a greater degree of 5 base pair mismatching than could occur under "highly stringent conditions" is able to form. Examples of typical "moderately stringent conditions" are 0.015M sodium chloride, 0.0015M sodium citrate at 50-65°C or 0.015M sodium chloride, 0.0015M sodium citrate, and 20% formamide at 37-50°C. By way of 10 example, a "moderately stringent" condition of 50°C in 0.015 M sodium ion will allow about a 21% mismatch.

It will be appreciated by those skilled in the art that there is no absolute distinction between "highly" and "moderately" stringent conditions. For example, at 0.015M 15 sodium ion (no formamide), the melting temperature of perfectly matched long DNA is about 71°C. With a wash at 65°C (at the same ionic strength), this would allow for approximately a 6% mismatch. To capture more distantly related sequences, one skilled in the art can simply lower the 20 temperature or raise the ionic strength.

A good estimate of the melting temperature in 1M NaCl* for oligonucleotide probes up to about 20nt is given by:

$$T_m = 2^\circ\text{C per A-T base pair} + 4^\circ\text{C per G-C base pair}$$

*The sodium ion concentration in 6X salt sodium citrate 25 (SSC) is 1M. See Suggs *et al.*, Developmental Biology Using Purified Genes, p. 683, Brown and Fox (eds.) (1981).

High stringency washing conditions for oligonucleotides are usually at a temperature of 0-5°C below the T_m of the oligonucleotide in 6X SSC, 0.1% SDS.

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In another embodiment, related nucleic acid molecules comprise or consist of a nucleotide sequence that is about 70 percent identical to the nucleotide sequence as shown in SEQ ID NO: 1, or comprise or consist essentially of a nucleotide sequence encoding a polypeptide that is about 70 percent identical to the polypeptide as set forth in SEQ ID NO: 2. In preferred embodiments, the nucleotide sequences are about 75 percent, or about 80 percent, or about 85 percent, or about 90 percent, or about 95, 96, 97, 98, or 99 percent identical to the nucleotide sequence as shown in SEQ ID NO: 1, or the nucleotide sequences encode a polypeptide that is about 75 percent, or about 80 percent, or about 85 percent, or about 90 percent, or about 95, 96, 97, 98, or 99 percent identical to the polypeptide sequence as set forth in SEQ ID NO: 2.

Differences in the nucleic acid sequence may result in conservative and/or non-conservative modifications of the amino acid sequence relative to the amino acid sequence of SEQ ID NO: 2.

Conservative modifications to the amino acid sequence of SEQ ID NO: 2 (and the corresponding modifications to the encoding nucleotides) will produce IL-17 receptor like polypeptides having functional and chemical characteristics similar to those of naturally occurring IL-17 receptor like polypeptide. In contrast, substantial modifications in the functional and/or chemical characteristics of IL-17 receptor like polypeptides may be accomplished by selecting substitutions in the amino acid sequence of SEQ ID NO: 2 that differ significantly in their effect on maintaining (a) the structure of the molecular backbone in the area of the substitution, for example, as a sheet or helical conformation, (b) the charge or hydrophobicity of the molecule at the target site, or (c) the bulk of the side chain.

For example, a "conservative amino acid substitution" may involve a substitution of a native amino acid residue with a

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nonnative residue such that there is little or no effect on the polarity or charge of the amino acid residue at that position. Furthermore, any native residue in the polypeptide may also be substituted with alanine, as has been previously 5 described for "alanine scanning mutagenesis."

Conservative amino acid substitutions also encompass non-naturally occurring amino acid residues which are typically incorporated by chemical peptide synthesis rather than by 10 synthesis in biological systems. These include peptidomimetics, and other reversed or inverted forms of amino acid moieties. It will be appreciated by those of skill in the art that nucleic acid polypeptide molecules described herein may be chemically synthesized as well as produced by 15 recombinant means.

Naturally occurring residues may be divided into classes based on common side chain properties:

- 1) hydrophobic: norleucine, Met, Ala, Val, Leu, Ile;
- 2) neutral hydrophilic: Cys, Ser, Thr, Asn, Gln;
- 3) acidic: Asp, Glu;
- 4) basic: His, Lys, Arg;
- 5) residues that influence chain orientation: Gly, Pro; and
- 6) aromatic: Trp, Tyr, Phe.

For example, non-conservative substitutions may involve 25 the exchange of a member of one of these classes for a member from another class. Such substituted residues may be introduced into regions of the human IL-17 receptor like polypeptide that are homologous with non-human IL-17 receptor 30 like polypeptide orthologs, or into the non-homologous regions of the molecule.

In making such changes, the hydropathic index of amino acids may be considered. Each amino acid has been assigned a

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hydropathic index on the basis of their hydrophobicity and charge characteristics, these are: isoleucine (+4.5); valine (+4.2); leucine (+3.8); phenylalanine (+2.8); cysteine/cystine (+2.5); methionine (+1.9); alanine (+1.8); glycine (-0.4); 5 threonine (-0.7); serine (-0.8); tryptophan (-0.9); tyrosine (-1.3); proline (-1.6); histidine (-3.2); glutamate (-3.5); glutamine (-3.5); aspartate (-3.5); asparagine (-3.5); lysine (-3.9); and arginine (-4.5).

The importance of the hydropathic amino acid index in 10 conferring interactive biological function on a protein is understood in the art. Kyte et al., *J. Mol. Biol.*, 157:105-131 (1982). It is known that certain amino acids may be substituted for other amino acids having a similar hydropathic index or score and still retain a similar biological activity. 15 In making changes based upon the hydropathic index, the substitution of amino acids whose hydropathic indices are within ± 2 is preferred, those which are within ± 1 are particularly preferred, and those within ± 0.5 are even more particularly preferred.

It is also understood in the art that the substitution of 20 like amino acids can be made effectively on the basis of hydrophilicity, particularly where the biologically functionally equivalent protein or peptide thereby created is intended for use in immunological embodiments, as in the 25 present case. The greatest local average hydrophilicity of a protein, as governed by the hydrophilicity of its adjacent amino acids, correlates with its immunogenicity and antigenicity, i.e., with a biological property of the protein.

The following hydrophilicity values have been assigned to 30 amino acid residues: arginine (+3.0); lysine (+3.0); aspartate (+3.0 \pm 1); glutamate (+3.0 \pm 1); serine (+0.3); asparagine (+0.2); glutamine (+0.2); glycine (0); threonine (-0.4); proline (-0.5 \pm 1); alanine (-0.5); histidine (-0.5); cysteine

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(-1.0); methionine (-1.3); valine (-1.5); leucine (-1.8); isoleucine (-1.8); tyrosine (-2.3); phenylalanine (-2.5); tryptophan (-3.4). In making changes based upon similar hydrophilicity values, the substitution of amino acids whose hydrophilicity values are within ± 2 is preferred, those which are within ± 1 are particularly preferred, and those within ± 0.5 are even more particularly preferred. One may also identify epitopes from primary amino acid sequences on the basis of hydrophilicity. These regions are also referred to 10 as "epitopic core regions."

Desired amino acid substitutions (whether conservative or non-conservative) can be determined by those skilled in the art at the time such substitutions are desired. For example, amino acid substitutions can be used to identify important 15 residues of the IL-17 receptor like polypeptide, or to increase or decrease the affinity of the IL-17 receptor like polypeptides described herein.

Exemplary amino acid substitutions are set forth in Table I.

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Table IAmino Acid Substitutions

Original Residues	Exemplary Substitutions	Preferred Substitutions
Ala	Val, Leu, Ile	Val
Arg	Lys, Gln, Asn	Lys
Asn	Gln	Gln
Asp	Glu	Glu
Cys	Ser, Ala	Ser
Gln	Asn	Asn
Glu	Asp	Asp
Gly	Pro, Ala	Ala
His	Asn, Gln, Lys, Arg	Arg
Ile	Leu, Val, Met, Ala, Phe, Norleucine	Leu
Leu	Norleucine, Ile, Val, Met, Ala, Phe	Ile
Lys	Arg, 1,4 Diamino- butyric Acid, Gln, Asn	Arg
Met	Leu, Phe, Ile	Leu
Phe	Leu, Val, Ile, Ala, Tyr	Leu
Pro	Ala	Gly
Ser	Thr, Ala, Cys	Thr
Thr	Ser	Ser
Trp	Tyr, Phe	Tyr
Tyr	Trp, Phe, Thr, Ser	Phe
Val	Ile, Met, Leu, Phe, Ala, Norleucine	Leu

A skilled artisan will be able to determine suitable
5 variants of the polypeptide as set forth in SEQ ID NO: 2 using
well known techniques. For identifying suitable areas of the

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molecule that may be changed without destroying biological activity. Also, one skilled in the art will realize that even areas that may be important for biological activity or for structure may be subject to conservative amino acid 5 substitutions without destroying the biological activity or without adversely affecting the polypeptide structure.

For example, when similar polypeptides with similar activities from the same species or from other species are known, one skilled in the art may compare the amino acid 10 sequence of an IL-17 receptor like polypeptide to such similar polypeptides. With such a comparison, one can identify residues and portions of the molecules that are conserved among similar polypeptides. It will be appreciated that changes in areas of an IL-17 receptor like polypeptide that 15 are not conserved relative to such similar polypeptides would be less likely to adversely affect the biological activity and/or structure of the IL-17 receptor like polypeptide. One skilled in the art would also know that, even in relatively conserved regions, one may substitute chemically similar amino 20 acids for the naturally occurring residues while retaining activity (conservative amino acid residue substitutions). Therefore, even areas that may be important for biological activity or for structure may be subject to conservative amino acid substitutions without destroying the biological activity 25 or without adversely affecting the polypeptide structure.

For predicting suitable areas of the molecule that may be changed without destroying activity, one skilled in the art may target areas not believed to be important for activity. For example, when similar polypeptides with similar activities 30 from the same species or from other species are known, one skilled in the art may compare the amino acid sequence of IL-17 receptor like polypeptide to such similar polypeptides. After making such a comparison, one skilled in the art can

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determine residues and portions of the molecules that are conserved among similar polypeptides. One skilled in the art would know that changes in areas of the IL-17 receptor like molecule that are not conserved would be less likely to 5 adversely affect the biological activity and/or structure of a IL-17 receptor like polypeptide. One skilled in the art would also know that, even in relatively conserved regions, one may substitute chemically similar amino acids for the naturally occurring residues while retaining activity (conservative 10 amino acid residue substitutions).

Additionally, one skilled in the art can review structure-function studies identifying residues in similar polypeptides that are important for activity or structure. In view of such a comparison, one can predict the importance of 15 amino acid residues in an IL-17 receptor like polypeptide that correspond to amino acid residues that are important for activity or structure in similar polypeptides. One skilled in the art may opt for chemically similar amino acid substitutions for such predicted important amino acid residues 20 of IL-17 receptor like polypeptides.

One skilled in the art can also analyze the three-dimensional structure and amino acid sequence in relation to that structure in similar polypeptides. In view of that information, one skilled in the art may predict the alignment 25 of amino acid residues of an IL-17 receptor like polypeptide with respect to its three dimensional structure. One skilled in the art may choose not to make radical changes to amino acid residues predicted to be on the surface of the protein, since such residues may be involved in important interactions 30 with other molecules. Moreover, one skilled in the art may generate test variants containing a single amino acid substitution at each desired amino acid residue. The variants can then be screened using activity assays known to those skilled in the art. Such variants could be used to gather

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information about suitable variants. For example, if one discovered that a change to a particular amino acid residue resulted in destroyed, undesirably reduced, or unsuitable activity, variants with such a change would be avoided. In 5 other words, based on information gathered from such routine experiments, one skilled in the art can readily determine the amino acids where further substitutions should be avoided either alone or in combination with other mutations.

A number of scientific publications have been devoted to 10 the prediction of secondary structure. See Moult J., *Curr. Op. in Biotech.*, 7(4):422-427 (1996), Chou et al., *Biochemistry*, 13(2):222-245 (1974); Chou et al., *Biochemistry*, 113(2):211-222 (1974); Chou et al., *Adv. Enzymol. Relat. Areas Mol. Biol.*, 47:45-148 (1978); Chou et al., *Ann. Rev. Biochem.*, 47:251-276 and Chou et al., *Biophys. J.*, 26:367-384 (1979). Moreover, computer programs are currently available to assist 15 with predicting secondary structure. One method of predicting secondary structure is based upon homology modeling. For example, two polypeptides or proteins which have a sequence identity of greater than 30%, or similarity greater than 40% 20 often have similar structural topologies. The recent growth of the protein structural data base (PDB) has provided enhanced predictability of secondary structure, including the potential number of folds within a polypeptide's or protein's 25 structure. See Holm et al., *Nucl. Acid. Res.*, 27(1):244-247 (1999). It has been suggested (Brenner et al., *Curr. Op. Struct. Biol.*, 7(3):369-376 (1997)) that there are a limited 30 number of folds in a given polypeptide or protein and that once a critical number of structures have been resolved, structural prediction will gain dramatically in accuracy.

Additional methods of predicting secondary structure include "threading" (Jones, D., *Curr. Opin. Struct. Biol.*, 7(3):377-87 (1997); Sippl et al., *Structure*, 4(1):15-9 (1996)), "profile analysis" (Bowie et al., *Science*, 253:164-

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170 (1991); Gribskov et al., *Meth. Enzym.*, 183:146-159 (1990);
Gribskov et al., *Proc. Nat. Acad. Sci.*, 84(13):4355-4358
(1987)), and "evolutionary linkage" (See Home, *supra*, and
Brenner, *supra*).

5 IL-17 receptor like polypeptide analogs of the invention
can be determined by comparing the amino acid sequence of IL-
17 receptor like polypeptide with related family members. An
exemplary IL-17 receptor like polypeptide related family
member is a human IL-17 receptor. This comparison can be
10 accomplished by using a Pileup alignment (Wisconsin GCG
Program Package) or an equivalent (overlapping) comparison
with multiple family members within conserved and non-
conserved regions.

As shown in Figure 2, the predicted amino acid sequences
15 of human IL-17 receptor like polypeptides (SEQ ID NO: 2) is
aligned with a known human IL-17 receptor family member (SEQ
ID NO: 3). Other IL-17 receptor like polypeptide analogs can
be determined using these or other methods known to those of
skill in the art. These overlapping sequences provide
20 guidance for conservative and non-conservative amino acids
substitutions resulting in additional IL-17 receptor like
analogs. It will be appreciated that these amino acid
substitutions can consist of naturally occurring or non-
naturally occurring amino acids. For example, potential IL-17
25 receptor like analogs may have the Gly at residue at position
45 of SEQ ID NO: 2 substituted with a Pro or Ala residue, the
Phe residue at position 227 of SEQ ID NO: 2 substituted with a
Leu, Val, Ile, Ala or Tyr residue, and/or the Ser residue at
position 363 of SEQ ID NO: 2 substituted with a Thr, Ala or
30 Cys. In addition, potential IL-17 receptor like analogs may
have the Val residue at position 374 of SEQ ID NO: 2
substituted with a Ile, Met, Leu, Phe, Ala or norleucine
residue, the Cys residue at position 385 of SEQ ID NO: 2,
substituted with a Ser or Ala residue, the Asp residue at
35 position 515 of SEQ ID NO: 2 substituted with a Glu residue

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and/or the Cys residue at position 602 substituted with a Ser or Ala.

Preferred IL-17 receptor like polypeptide variants include glycosylation variants wherein the number and/or type 5 of glycosylation sites has been altered compared to the amino acid sequence set forth in SEQ ID NO: 2. In one embodiment, IL-17 receptor like polypeptide variants comprise a greater or a lesser number of N-linked glycosylation sites than the amino acid sequence set forth in SEQ ID NO: 2. An N-linked 10 glycosylation site is characterized by the sequence: Asn-X-Ser or Asn-X-Thr, wherein the amino acid residue designated as X may be any amino acid residue except proline. The substitution(s) of amino acid residues to create this sequence provides a potential new site for the addition of an N-linked 15 carbohydrate chain. Alternatively, substitutions which eliminate this sequence will remove an existing N-linked carbohydrate chain. Also provided is a rearrangement of N-linked carbohydrate chains wherein one or more N-linked glycosylation sites (typically those that are naturally 20 occurring) are eliminated and one or more new N-linked sites are created. Additional preferred IL-17 receptor like variants include cysteine variants, wherein one or more cysteine residues are deleted from or substituted for another amino acid (e.g., serine) as compared to the amino acid 25 sequence set forth in SEQ ID NO: 2. Cysteine variants are useful when IL-17 receptor like polypeptides must be refolded into a biologically active conformation such as after the isolation of insoluble inclusion bodies. Cysteine variants generally have fewer cysteine residues than the native 30 protein, and typically have an even number to minimize interactions resulting from unpaired cysteines.

In addition, the polypeptide comprising the amino acid sequence of SEQ ID NO: 2 or an IL-17 receptor like polypeptide variant may be fused to a homologous polypeptide to form a

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homodimer or to a heterologous polypeptide to form a heterodimer. Heterologous peptides and polypeptides include, but are not limited to: an epitope to allow for the detection and/or isolation of an IL-17 receptor like fusion polypeptide; 5 a transmembrane receptor protein or a portion thereof, such as an extracellular domain, or a transmembrane and intracellular domain; a ligand or a portion thereof which binds to a transmembrane receptor protein; an enzyme or portion thereof which is catalytically active; a polypeptide or peptide which promotes oligomerization, such as a leucine zipper domain; a 10 polypeptide or peptide which increases stability, such as an immunoglobulin constant region; and a polypeptide which has a therapeutic activity different from the polypeptide comprising the amino acid sequence as set forth in SEQ ID NO: 2 or an IL- 15 17 receptor like polypeptide variant. In addition, a IL-17 receptor like polypeptide may be fused to itself or to a fragment, variant, or derivative thereof.

Fusions can be made either at the amino terminus or at the carboxy terminus of the polypeptide comprising the amino 20 acid sequence set forth in SEQ ID NO: 2 or an IL-17 receptor like polypeptide variant. Fusions may be direct with no linker or adapter molecule or indirect using a linker or adapter molecule. A linker or adapter molecule may be one or more amino acid residues, typically up to about 20 amino acids 25 residues, or up to about 50 amino acid residues. A linker or adapter molecule may also be designed with a cleavage site for a DNA restriction endonuclease or for a protease to allow for the separation of the fused moieties. It will be appreciated that once constructed, the fusion polypeptides can be 30 derivatized according to the methods described herein.

In a further embodiment of the invention, the polypeptide comprising the amino acid sequence of SEQ ID NO: 2 or an IL-17 receptor like polypeptide variant, including a fragment, variant and/or derivative, is fused to one or more domains of

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an Fc region of human IgG. Antibodies comprise two functionally independent parts, a variable domain known as "Fab", which binds antigen, and a constant domain known as "Fc", which is involved in effector functions such as 5 complement activation and attack by phagocytic cells. An Fc has a long serum half-life, whereas an Fab is short-lived. Capon et al., *Nature*, 337:525-31 (1989). When constructed together with a therapeutic protein, an Fc domain can provide longer half-life or incorporate such functions as Fc receptor 10 binding, protein A binding, complement fixation and perhaps even placental transfer. *Id.* Table II summarizes the use of certain Fc fusions known in the art, including materials and methods applicable to the production of fused IL-17 receptor like polypeptide.

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Table II
Fc Fusion with Therapeutic Proteins

Form of Fc	Fusion partner	Therapeutic implications	Reference
IgG1	N-terminus of CD30-L	Hodgkin's disease; anaplastic lymphoma; T-cell leukemia	U.S. Patent No. 5,480,981
Murine Fc γ 2a	IL-10	anti-inflammatory; transplant rejection	Zheng et al. (1995), <i>J. Immunol.</i> , <u>154</u> : 5590-5600
IgG1	TNF receptor	septic shock	Fisher et al. (1996), <i>N. Engl. J. Med.</i> , <u>334</u> : 1697-1702; Van Zee et al., (1996), <i>J. Immunol.</i> , <u>156</u> : 2221-2230
IgG, IgA, IgM, or IgE (excluding the first domain)	TNF receptor	inflammation, autoimmune disorders	U.S. Pat. No. 5,808,029, issued September 15, 1998
IgG1	CD4 receptor	AIDS	Capon et al. (1989), <i>Nature</i> <u>337</u> : 525-531
IgG1, IgG3	N-terminus of IL-2	anti-cancer, antiviral	Harvill et al. (1995), <i>Immunotech.</i> , <u>1</u> : 95-105
IgG1	C-terminus of OPG	osteoarthritis; bone density	WO 97/23614, published July 3, 1997
IgG1	N-terminus of leptin	anti-obesity	PCT/US 97/23183, filed December 11, 1997
Human Ig C γ 1	CTLA-4	autoimmune disorders	Linsley (1991), <i>J. Exp. Med.</i> , <u>174</u> :561-569

In one example, all or a portion of the human IgG hinge,
5 CH2 and CH3 regions may be fused at either the N-terminus or C-terminus of the IL-17 receptor like polypeptides using

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methods known to the skilled artisan. In another example, a portion of hinge regions and CH₂ and CH₃ regions may be fused. The resulting IL-17 receptor like fusion polypeptide may be purified by use of a Protein A affinity column. Peptides and 5 proteins fused to an Fc region have been found to exhibit a substantially greater half-life *in vivo* than the unfused counterpart. Also, a fusion to an Fc region allows for dimerization/multimerization of the fusion polypeptide. The Fc region may be a naturally occurring Fc region, or may be 10 altered to improve certain qualities, such as therapeutic qualities, circulation time, reduce aggregation, etc.

Identity and similarity of related nucleic acid molecules and polypeptides can be readily calculated by known methods. Such methods include, but are not limited to, those described 15 in Computational Molecular Biology, Lesk, A.M., ed., Oxford University Press, New York, 1988; Biocomputing: Informatics and Genome Projects, Smith, D.W., ed., Academic Press, New York, 1993; Computer Analysis of Sequence Data, Part 1, Griffin, A.M., and Griffin, H.G., eds., Humana Press, New Jersey, 1994; Sequence Analysis in Molecular Biology, von Heinje, G., Academic Press, 1987; Sequence Analysis Primer, Gribskov, M. and Devereux, J., eds., M. Stockton Press, New York, 1991; and Carillo et al., *SIAM J. Applied Math.*, 48:1073 (1988).

25 Preferred methods to determine identity and/or similarity are designed to give the largest match between the sequences tested. Methods to determine identity and similarity are described in publicly available computer programs. Preferred computer program methods to determine identity and similarity 30 between two sequences include, but are not limited to, the GCG program package, including GAP (Devereux et al., *Nucl. Acid. Res.*, 12:387 (1984); Genetics Computer Group, University of Wisconsin, Madison, WI), BLASTP, BLASTN, and FASTA (Altschul et al., *J. Mol. Biol.*, 215:403-410 (1990)). The BLASTX

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program is publicly available from the National Center for Biotechnology Information (NCBI) and other sources (*BLAST Manual*, Altschul et al. NCB/NLM/NIH Bethesda, MD 20894; Altschul et al., *supra*). The well known Smith Waterman 5 algorithm may also be used to determine identity.

Certain alignment schemes for aligning two amino acid sequences may result in the matching of only a short region of the two sequences, and this small aligned region may have very high sequence identity even though there is no significant 10 relationship between the two full length sequences. Accordingly, in a preferred embodiment, the selected alignment method (GAP program) will result in an alignment that spans at least 50 contiguous amino acids of the target polypeptide.

For example, using the computer algorithm GAP (Genetics 15 Computer Group, University of Wisconsin, Madison, WI), two polypeptides for which the percent sequence identity is to be determined are aligned for optimal matching of their respective amino acids (the "matched span", as determined by the algorithm). A gap opening penalty (which is calculated as 20 3X the average diagonal; the "average diagonal" is the average of the diagonal of the comparison matrix being used; the "diagonal" is the score or number assigned to each perfect amino acid match by the particular comparison matrix) and a gap extension penalty (which is usually 1/10 times the gap 25 opening penalty), as well as a comparison matrix such as PAM 250 or BLOSUM 62 are used in conjunction with the algorithm. A standard comparison matrix (see Dayhoff et al., Atlas of Protein Sequence and Structure, vol. 5, supp.3 (1978) for the PAM 250 comparison matrix; Henikoff et al., Proc. 30 Natl. Acad. Sci USA, 89:10915-10919 (1992) for the BLOSUM 62 comparison matrix) is also used by the algorithm.

Preferred parameters for a polypeptide sequence comparison include the following:

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Algorithm: Needleman et al., *J. Mol. Biol.*, 48:443-453 (1970);

5. Comparison matrix: BLOSUM 62 from Henikoff et al., *Proc. Natl. Acad. Sci. USA*, 89:10915-10919 (1992);

Gap Penalty: 12

Gap Length Penalty: 4

Threshold of Similarity: 0

10 The GAP program is useful with the above parameters. The aforementioned parameters are the default parameters for polypeptide comparisons (along with no penalty for end gaps) using the GAP algorithm.

15 Preferred parameters for nucleic acid molecule sequence comparisons include the following:

Algorithm: Needleman et al., *J. Mol Biol.*, 48:443-453 (1970);

Comparison matrix: matches = +10, mismatch = 0

Gap Penalty: 50

20 Gap Length Penalty: 3

The GAP program is also useful with the above parameters. The aforementioned parameters are the default parameters for nucleic acid molecule comparisons.

25 Other exemplary algorithms, gap opening penalties, gap extension penalties, comparison matrices, thresholds of similarity, etc. may be used,, including those set forth in the Program Manual, Wisconsin Package, Version 9, September, 1997. The particular choices to be made will be apparent to 30 those of skill in the art and will depend on the specific comparison to be made, such as DNA to DNA, protein to protein, protein to DNA; and additionally, whether the comparison is

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between given pairs of sequences (in which case GAP or BestFit are generally preferred) or between one sequence and a large database of sequences (in which case FASTA or BLASTA are preferred).

5

Synthesis

It will be appreciated by those skilled in the art the nucleic acid and polypeptide molecules described herein may be produced by recombinant and other means.

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Nucleic Acid Molecules

The nucleic acid molecules encode a polypeptide comprising the amino acid sequence of an IL-17 receptor like polypeptide can readily be obtained in a variety of ways 15 including, without limitation, chemical synthesis, cDNA or genomic library screening, expression library screening and/or PCR amplification of cDNA.

Recombinant DNA methods used herein are generally those set forth in Sambrook et al., *Molecular Cloning: A Laboratory 20 Manual*, Cold Spring Harbor Laboratory Press, Cold Spring Harbor, NY (1989), and/or Ausubel et al., eds., *Current Protocols in Molecular Biology*, Green Publishers Inc. and Wiley and Sons, NY (1994). The present invention provides for nucleic acid molecules as described herein and methods for 25 obtaining the molecules.

A gene or cDNA encoding a IL-17 receptor like polypeptide or fragment thereof may be obtained by hybridization screening of a genomic or cDNA library , or by PCR amplification. Where a gene encoding the amino acid sequence of an IL-17 receptor 30 like polypeptide has been identified from one species, all or a portion of that gene may be used as a probe to identify orthologs or related genes from the same species. The probes or primers may be used to screen cDNA libraries from various

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tissue sources believed to express the IL-17 receptor like polypeptide. In addition, part or all of a nucleic acid molecule having the sequence as set forth in SEQ ID NO: 1 may be used to screen a genomic library to identify and isolate a 5 gene encoding the amino acid sequence of an IL-17 receptor like polypeptide. Typically, conditions of moderate or high stringency will be employed for screening to minimize the number of false positives obtained from the screen.

Nucleic acid molecules encoding the amino acid sequence 10 of IL-17 receptor like polypeptides may also be identified by expression cloning which employs the detection of positive clones based upon a property of the expressed protein. Typically, nucleic acid libraries are screened by the binding 15 of an antibody or other binding partner (e.g., receptor or ligand) to cloned proteins which are expressed and displayed on a host cell surface. The antibody or binding partner is modified with a detectable label to identify those cells expressing the desired clone.

Recombinant expression techniques conducted in accordance 20 with the descriptions set forth below may be followed to produce these polynucleotides and to express the encoded polypeptides. For example, by inserting a nucleic acid sequence which encodes the amino acid sequence of an IL-17 receptor like polypeptide into an appropriate vector, one 25 skilled in the art can readily produce large quantities of the desired nucleotide sequence. The sequences can then be used to generate detection probes or amplification primers. Alternatively, a polynucleotide encoding the amino acid sequence of an IL-17 receptor like polypeptide can be inserted 30 into an expression vector. By introducing the expression vector into an appropriate host, the encoded IL-17 receptor like polypeptide may be produced in large amounts.

Another method for obtaining a suitable nucleic acid sequence is the polymerase chain reaction (PCR). In this

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method, cDNA is prepared from poly(A)+RNA or total RNA using the enzyme reverse transcriptase. Two primers, typically complementary to two separate regions of cDNA (oligonucleotides) encoding the amino acid sequence of an IL-17 receptor like polypeptide, are then added to the cDNA along with a polymerase such as Tag polymerase, and the polymerase amplifies the cDNA region between the two primers.

Another means of preparing a nucleic acid molecule encoding the amino acid sequence of an IL-17 receptor like polypeptide, including a fragment or variant, is chemical synthesis using methods well known to the skilled artisan such as those described by Engels et al., *Angew. Chem. Intl. Ed.*, 28:716-734 (1989). These methods include, *inter alia*, the phosphotriester, phosphoramidite, and H-phosphonate methods for nucleic acid synthesis. A preferred method for such chemical synthesis is polymer-supported synthesis using standard phosphoramidite chemistry. Typically, the DNA encoding the amino acid sequence of an IL-17 receptor like polypeptide will be several hundred nucleotides in length. Nucleic acids larger than about 100 nucleotides can be synthesized as several fragments using these methods. The fragments can then be ligated together to form the full length nucleotide sequence of an IL-17 receptor like polypeptide. Usually, the DNA fragment encoding the amino terminus of the polypeptide will have an ATG, which encodes a methionine residue. This methionine may or may not be present on the mature form of the IL-17 receptor like polypeptide, depending on whether the polypeptide produced in the host cell is designed to be secreted from that cell. Other methods known to the skilled artisan may be used as well.

In some cases, it may be desirable to prepare nucleic acid molecules encoding IL-17 receptor like polypeptide variants. Nucleic acid molecules encoding variants may be produced using site directed mutagenesis, PCR amplification, or other appropriate methods, where the primer(s) have the desired point

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mutations (see Sambrook et al., *supra*, and Ausubel et al., *supra*, for descriptions of mutagenesis techniques). Chemical synthesis using methods described by Engels et al., *supra*, may also be used to prepare such variants. Other methods known to
5 the skilled artisan may be used as well.

In certain embodiments, nucleic acid variants contain codons which have been altered for the optimal expression of an IL-17 receptor like polypeptide in a given host cell. Particular codon alterations will depend upon the IL-17
10 receptor like polypeptide(s) and host cell(s) selected for expression. Such "codon optimization" can be carried out by a variety of methods, for example, by selecting codons which are preferred for use in highly expressed genes in a given host cell. Computer algorithms which incorporate codon frequency
15 tables such as "Ecohigh.cod" for codon preference of highly expressed bacterial genes may be used and are provided by the University of Wisconsin Package Version 9.0, Genetics Computer Group, Madison, WI. Other useful codon frequency tables include "Celegans_high.cod", "Celegans_low.cod",
20 "Drosophila_high.cod", "Human_high.cod", "Maize_high.cod", and "Yeast_high.cod".

In other embodiments, nucleic acid molecules encode IL-17 receptor like variants with conservative amino acid substitutions as described herein, IL-17 receptor like variants comprising an addition and/or a deletion of one or more N-linked or O-linked glycosylation sites, IL-17 receptor like variants having deletions and/or substitutions of one or more cysteine residues, or IL-17 receptor like polypeptide fragments as described herein. In addition, nucleic acid molecules may
25 encode any combination of IL-17 receptor like variants, fragments, and fusion polypeptides described herein.

Vectors and Host Cells

A nucleic acid molecule encoding the amino acid sequence of an IL-17 receptor like polypeptide may be inserted into an

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appropriate expression vector using standard ligation techniques. The vector is typically selected to be functional in the particular host cell employed (i.e., the vector is compatible with the host cell machinery such that amplification of the gene and/or expression of the gene can occur). A nucleic acid molecule encoding the amino acid sequence of an IL-17 receptor like polypeptide may be amplified/expressed in prokaryotic, yeast, insect (baculovirus systems), and/or eukaryotic host cells. Selection of the host cell will depend in part on whether an IL-17 receptor like polypeptide is to be post-translationally modified (e.g., glycosylated and/or phosphorylated). If so; yeast, insect, or mammalian host cells are preferable. For a review of expression vectors, see *Meth. Enz.*, v.185, D.V. Goeddel, ed. Academic Press Inc., San Diego, CA (1990).

Typically, expression vectors used in any of the host cells will contain sequences for plasmid maintenance and for cloning and expression of exogenous nucleotide sequences. Such sequences, collectively referred to as "flanking sequences" in certain embodiments will typically include one or more of the following nucleotide sequences: a promoter, one or more enhancer sequences, an origin of replication, a transcriptional termination sequence, a complete intron sequence containing a donor and acceptor splice site, a sequence encoding a leader sequence for polypeptide secretion, a ribosome binding site, a polyadenylation sequence, a polylinker region for inserting the nucleic acid encoding the polypeptide to be expressed, and a selectable marker element. Each of these sequences is discussed below.

Optionally, the vector may contain a "tag"-encoding sequence, i.e., an oligonucleotide molecule located at the 5' or 3' end of the IL-17 receptor like polypeptide coding sequence; the oligonucleotide sequence encodes polyHis (such as hexaHis), or other "tag" such as FLAG, HA (hemagglutinin Influenza virus) or myc for which commercially available

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antibodies exist. This tag is typically fused to the polypeptide upon expression of the polypeptide, and can serve as a means for affinity purification of the IL-17 receptor like polypeptide from the host cell. Affinity purification
5 can be accomplished, for example, by column chromatography using antibodies against the tag as an affinity matrix. Optionally, the tag can subsequently be removed from the purified IL-17 receptor like polypeptide by various means such as using certain peptidases for cleavage.

10 Flanking sequences may be homologous (i.e., from the same species and/or strain as the host cell), heterologous (i.e., from a species other than the host cell species or strain), hybrid (i.e., a combination of flanking sequences from more than one source) or synthetic, or the flanking sequences may
15 be native sequences which normally function to regulate IL-17 receptor like polypeptide expression. As such, the source of a flanking sequence may be any prokaryotic or eukaryotic organism, any vertebrate or invertebrate organism, or any plant, provided that the flanking sequence is functional in,
20 and can be activated by, the host cell machinery.

The flanking sequences useful in the vectors of this invention may be obtained by any of several methods well known in the art. Typically, flanking sequences useful herein other than the IL-17 receptor like gene flanking sequences will have
25 been previously identified by mapping and/or by restriction endonuclease digestion and can thus be isolated from the proper tissue source using the appropriate restriction endonucleases. In some cases, the full nucleotide sequence of a flanking sequence may be known. Here, the flanking sequence
30 may be synthesized using the methods described herein for nucleic acid synthesis or cloning.

Where all or only a portion of the flanking sequence is known, it may be obtained using PCR and/or by screening a genomic library with suitable oligonucleotide and/or flanking

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sequence fragments from the same or another species. Where the flanking sequence is not known, a fragment of DNA containing a flanking sequence may be isolated from a larger piece of DNA that may contain, for example, a coding sequence 5 or even another gene or genes. Isolation may be accomplished by restriction endonuclease digestion to produce the proper DNA fragment followed by isolation using agarose gel purification, Qiagen® column chromatography (Chatsworth, CA), or other methods known to the skilled artisan. The selection 10 of suitable enzymes to accomplish this purpose will be readily apparent to one of ordinary skill in the art.

An origin of replication is typically a part of those prokaryotic expression vectors purchased commercially, and the origin aids in the amplification of the vector in a host cell. 15 Amplification of the vector to a certain copy number can, in some cases, be important for the optimal expression of an IL-17 receptor like polypeptide. If the vector of choice does not contain an origin of replication site, one may be chemically synthesized based on a known sequence, and ligated 20 into the vector. For example, the origin of replication from the plasmid pBR322 (Product No. 303-3s, New England Biolabs, Beverly, MA) is suitable for most Gram-negative bacteria and various origins (e.g., SV40, polyoma, adenovirus, vesicular stomatitis virus (VSV) or papillomaviruses such as HPV or BPV) 25 are useful for cloning vectors in mammalian cells. Generally, the origin of replication component is not needed for mammalian expression vectors (for example, the SV40 origin is often used only because it contains the early promoter).

A transcription termination sequence is typically located 30 3' of the end of a polypeptide coding region and serves to terminate transcription. Usually, a transcription termination sequence in prokaryotic cells is a G-C rich fragment followed by a poly T sequence. While the sequence is easily cloned from a library or even purchased commercially as part of a

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vector, it can also be readily synthesized using methods for nucleic acid synthesis such as those described herein.

A selectable marker gene element encodes a protein necessary for the survival and growth of a host cell grown in 5 a selective culture medium. Typical selection marker genes encode proteins that (a) confer resistance to antibiotics or other toxins, e.g., ampicillin, tetracycline, or kanamycin for prokaryotic host cells, (b) complement auxotrophic deficiencies of the cell; or (c) supply critical nutrients not 10 available from complex media. Preferred selectable markers are the kanamycin resistance gene, the ampicillin resistance gene, and the tetracycline resistance gene. A neomycin resistance gene may also be used for selection in prokaryotic and eukaryotic host cells.

15 Other selection genes may be used to amplify the gene which will be expressed. Amplification is the process wherein genes which are in greater demand for the production of a protein critical for growth are reiterated in tandem within the chromosomes of successive generations of recombinant 20 cells. Examples of suitable selectable markers for mammalian cells include dihydrofolate reductase (DHFR) and thymidine kinase. The mammalian cell transformants are placed under selection pressure which only the transformants are uniquely adapted to survive by virtue of the selection gene present in 25 the vector. Selection pressure is imposed by culturing the transformed cells under conditions in which the concentration of selection agent in the medium is successively changed, thereby leading to the amplification of both the selection gene and the DNA that encodes an IL-17 receptor like 30 polypeptide. As a result, increased quantities of IL-17 receptor like polypeptide are synthesized from the amplified DNA.

A ribosome binding site is usually necessary for translation initiation of mRNA and is characterized by a

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Shine-Dalgarno sequence (prokaryotes) or a Kozak sequence (eukaryotes). The element is typically located 3' to the promoter and 5' to the coding sequence of an IL-17 receptor like polypeptide to be expressed. The Shine-Dalgarno sequence 5 is varied but is typically a polypurine (i.e., having a high A-G content). Many Shine-Dalgarno sequences have been identified, each of which can be readily synthesized using methods set forth herein and used in a prokaryotic vector.

A leader, or signal, sequence may be used to direct an 10 IL-17 receptor like polypeptide out of the host cell. Typically, a nucleotide sequence encoding the signal sequence is positioned in the coding region of an IL-17 receptor like nucleic acid molecule, or directly at the 5' end of an IL-17 receptor like polypeptide coding region. Many signal 15 sequences have been identified, and any of those that are functional in the selected host cell may be used in conjunction with an IL-17 receptor like nucleic acid molecule. Therefore, a signal sequence may be homologous (naturally occurring) or heterologous to an IL-17 receptor like gene or 20 cDNA. Additionally, a signal sequence may be chemically synthesized using methods described herein. In most cases, the secretion of an IL-17 receptor like polypeptide from the host cell via the presence of a signal peptide will result in the removal of the signal peptide from the secreted IL-17 25 receptor like polypeptide. The signal sequence may be a component of the vector, or it may be a part of an IL-17 receptor like nucleic acid molecule that is inserted into the vector.

Included within the scope of this invention is the use of 30 either a nucleotide sequence encoding a native IL-17 receptor like polypeptide signal sequence joined to an IL-17 receptor like polypeptide coding region or a nucleotide sequence encoding a heterologous signal sequence joined to an IL-17 receptor like polypeptide coding region. The heterologous 35 signal sequence selected should be one that is recognized and

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processed, i.e., cleaved by a signal peptidase, by the host cell. For prokaryotic host cells that do not recognize and process the native IL-17 receptor like polypeptide signal sequence, the signal sequence is substituted by a prokaryotic signal sequence selected, for example, from the group of the alkaline phosphatase, penicillinase, or heat-stable enterotoxin II leaders. For yeast secretion, the native IL-17 receptor like polypeptide signal sequence may be substituted by the yeast invertase, alpha factor, or acid phosphatase leaders. In mammalian cell expression the native signal sequence is satisfactory, although other mammalian signal sequences may be suitable.

In some cases, such as where glycosylation is desired in a eukaryotic host cell expression system, one may manipulate the various presequences to improve glycosylation or yield. For example, one may alter the peptidase cleavage site of a particular signal peptide, or add presequences, which also may affect glycosylation. The final protein product may have, in the -1 position (relative to the first amino acid of the mature protein) one or more additional amino acids incident to expression, which may not have been totally removed. For example, the final protein product may have one or two amino acid residues found in the peptidase cleavage site, attached to the N-terminus. Alternatively, use of some enzyme cleavage sites may result in a slightly truncated form of the desired IL-17 receptor like polypeptide, if the enzyme cuts at such area within the mature polypeptide.

In many cases, transcription of a nucleic acid molecule is increased by the presence of one or more introns in the vector; this is particularly true where a polypeptide is produced in eukaryotic host cells, especially mammalian host cells. The introns used may be naturally occurring within the IL-17 receptor like gene, especially where the gene used is a full length genomic sequence or a fragment thereof. Where the intron is not naturally occurring within the gene (as for most

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cDNAs), the intron(s) may be obtained from another source. The position of the intron with respect to flanking sequences and the IL-17 receptor like gene is generally important, as the intron must be transcribed to be effective. Thus, when an 5 IL-17 receptor like cDNA molecule is being transcribed, the preferred position for the intron is 3' to the transcription start site, and 5' to the polyA transcription termination sequence. Preferably, the intron or introns will be located on one side or the other (i.e., 5' or 3') of the cDNA such 10 that it does not interrupt the coding sequence. Any intron from any source, including any viral, prokaryotic and eukaryotic (plant or animal) organisms, may be used to practice this invention, provided that it is compatible with the host cell(s) into which it is inserted. Also included 15 herein are synthetic introns. Optionally, more than one intron may be used in the vector.

The expression and cloning vectors of the present invention will each typically contain a promoter that is recognized by the host organism and operably linked to the 20 molecule encoding a IL-17 receptor like polypeptide. Promoters are untranscribed sequences located upstream (5') to the start codon of a structural gene (generally within about 100 to 1000 bp) that control the transcription of the structural gene. Promoters are conventionally grouped into 25 one of two classes, inducible promoters and constitutive promoters. Inducible promoters initiate increased levels of transcription from DNA under their control in response to some change in culture conditions, such as the presence or absence of a nutrient or a change in temperature. Constitutive 30 promoters, on the other hand, initiate continual gene product production; that is, there is little or no control over gene expression. A large number of promoters, recognized by a variety of potential host cells, are well known. A suitable promoter is operably linked to the DNA encoding an IL-17 35 receptor like polypeptide by removing the promoter from the

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source DNA by restriction enzyme digestion and inserting the desired promoter sequence into the vector. The native IL-17 receptor like gene promoter sequence may be used to direct amplification and/or expression of an IL-17 receptor like nucleic acid molecule. A heterologous promoter is preferred, however, if it permits greater transcription and higher yields of the expressed protein as compared to the native promoter, and if it is compatible with the host cell system that has been selected for use.

Promoters suitable for use with prokaryotic hosts include the beta-lactamase and lactose promoter systems; alkaline phosphatase, a tryptophan (trp) promoter system; and hybrid promoters such as the tac promoter. Other known bacterial promoters are also suitable. Their sequences have been published, thereby enabling one skilled in the art to ligate them to the desired DNA sequence(s), using linkers or adapters as needed to supply any useful restriction sites.

Suitable promoters for use with yeast hosts are also well known in the art. Yeast enhancers are advantageously used with yeast promoters. Suitable promoters for use with mammalian host cells are well known and include, but are not limited to, those obtained from the genomes of viruses such as polyoma virus, fowlpox virus, adenovirus (such as Adenovirus 2), bovine papilloma virus, avian sarcoma virus, cytomegalovirus (CMV), a retrovirus, hepatitis-B virus and most preferably Simian Virus 40 (SV40). Other suitable mammalian promoters include heterologous mammalian promoters, e.g., heat-shock promoters and the actin promoter.

Additional promoters which may be of interest in controlling IL-17 receptor like gene transcription include, but are not limited to: the SV40 early promoter region (Bernoist and Chambon, *Nature*, 290:304-310, 1981); the CMV promoter; the promoter contained in the 3' long terminal repeat of Rous sarcoma virus (Yamamoto et al., *Cell*, 22:787-

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797, 1980); the herpes thymidine kinase promoter (Wagner et al., *Proc. Natl. Acad. Sci. USA*, 78:144-1445, 1981); the regulatory sequences of the metallothioneine gene (Brinster et al., *Nature*, 296:39-42, 1982); prokaryotic expression vectors such as the beta-lactamase promoter (Villa-Kamaroff, et al., *Proc. Natl. Acad. Sci. USA*, 75:3727-3731, 1978); or the tac promoter (DeBoer, et al., *Proc. Natl. Acad. Sci. USA*, 80:21-25, 1983). Also of interest are the following animal transcriptional control regions, which exhibit tissue specificity and have been utilized in transgenic animals: the elastase I gene control region which is active in pancreatic acinar cells (Swift et al., *Cell*, 38:639-646, 1984; Ornitz et al., *Cold Spring Harbor Symp. Quant. Biol.*, 50:399-409 (1986); MacDonald, *Hepatology*, 7:425-515, 1987); the insulin gene control region which is active in pancreatic beta cells (Hanahan, *Nature*, 315:115-122, 1985); the immunoglobulin gene control region which is active in lymphoid cells (Grosschedl et al., *Cell*, 38:647-658 (1984); Adames et al., *Nature*, 318:533-538 (1985); Alexander et al., *Mol. Cell. Biol.*, 7:1436-1444, 1987); the mouse mammary tumor virus control region which is active in testicular, breast, lymphoid and mast cells (Leder et al., *Cell*, 45:485-495, 1986); the albumin gene control region which is active in liver (Pinkert et al., *Genes and Devel.*, 1:268-276, 1987); the alphafetoprotein gene control region which is active in liver (Krumlauf et al., *Mol. Cell. Biol.*, 5:1639-1648, 1985; Hammer et al., *Science*, 235:53-58, 1987); the alpha 1-antitrypsin gene control region which is active in the liver (Kelsey et al., *Genes and Devel.*, 1:161-171, 1987); the beta-globin gene control region which is active in myeloid cells (Mogram et al., *Nature*, 315:338-340, 1985; Kollias et al., *Cell*, 46:89-94, 1986); the myelin basic protein gene control region which is active in oligodendrocyte cells in the brain (Readhead et al., *Cell*, 48:703-712, 1987); the myosin light chain-2 gene control region which is active in skeletal muscle (Sani, *Nature*, 314:283-286, 1985); and the

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gonadotropic releasing hormone gene control region which is active in the hypothalamus (Mason *et al.*, *Science*, 234:1372-1378, 1986).

An enhancer sequence may be inserted into the vector to
5 increase the transcription of a DNA encoding an IL-17 receptor
like polypeptide of the present invention by higher
eukaryotes. Enhancers are *cis*-acting elements of DNA, usually
about 10-300 bp in length, that act on the promoter to
increase transcription. Enhancers are relatively orientation
10 and position independent. They have been found 5' and 3' to
the transcription unit. Several enhancer sequences available
from mammalian genes are known (e.g., globin, elastase,
albumin, alpha-feto-protein and insulin). Typically, however,
an enhancer from a virus will be used. The SV40 enhancer, the
15 cytomegalovirus early promoter enhancer, the polyoma enhancer,
and adenovirus enhancers are exemplary enhancing elements for
the activation of eukaryotic promoters. While an enhancer may
be spliced into the vector at a position 5' or 3' to an IL-17
RECEPTOR LIKE nucleic acid molecule, it is typically located
20 at a site 5' from the promoter.

Expression vectors of the invention may be constructed
from a starting vector such as a commercially available
vector. Such vectors may or may not contain all of the
desired flanking sequences. Where one or more of the desired
25 flanking sequences are not already present in the vector, they
may be individually obtained and ligated into the vector.
Methods used for obtaining each of the flanking sequences are
well known to one skilled in the art.

Preferred vectors for practicing this invention are those
30 which are compatible with bacterial, insect, and mammalian
host cells. Such vectors include, *inter alia*, pCRII, pCR3,
and pcDNA3.1 (Invitrogen Company, Carlsbad, CA), pBSII
(Stratagene Company, La Jolla, CA), pET15 (Novagen, Madison,
WI), pGEX (Pharmacia Biotech, Piscataway, NJ), pEGFP-N2

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(Clontech, Palo Alto, CA), pETL (BlueBacII; Invitrogen), pDSR-alpha (PCT Publication No. WO90/14363) and pFastBacDual (Gibco/BRL, Grand Island, NY).

Additional suitable vectors include, but are not limited
5 to, cosmids, plasmids or modified viruses, but it will be
appreciated that the vector system must be compatible with the
selected host cell. Such vectors include, but are not limited
to plasmids such as Bluescript[®] plasmid derivatives (a high
copy number ColE1-based phagemid, Stratagene Cloning Systems
10 Inc., La Jolla CA), PCR cloning plasmids designed for cloning
Taq-amplified PCR products (e.g., TOPO[™] TA Cloning[®] Kit,
PCR2.1[®] plasmid derivatives, Invitrogen, Carlsbad, CA), and
mammalian, yeast, or virus vectors such as a baculovirus
expression system (pBacPAK plasmid derivatives, Clontech, Palo
15 Alto, CA). The recombinant molecules can be introduced into
host cells via transformation, transfection, infection,
electroporation or other known techniques.

After the vector has been constructed and a nucleic acid
molecule encoding an IL-17 receptor like polypeptide has been
20 inserted into the proper site of the vector, the completed
vector may be inserted into a suitable host cell for
amplification and/or polypeptide expression. The
transformation of an expression vector for an IL-17 receptor
like polypeptide into a selected host cell may be accomplished
25 by well known methods including methods such as transfection,
infection, calcium chloride, electroporation, microinjection,
lipofection or the DEAE-dextran method or other known
techniques. The method selected will in part be a function of
the type of host cell to be used. These methods and other
30 suitable methods are well known to the skilled artisan, and
are set forth, for example, in Sambrook et al., *supra*.

Host cells may be prokaryotic host cells (such as *E. coli*) or eukaryotic host cells (such as a yeast cell, an
insect cell or a vertebrate cell). The host cell, when

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cultured under appropriate conditions, synthesizes an IL-17 receptor like polypeptide which can subsequently be collected from the culture medium (if the host cell secretes it into the medium) or directly from the host cell producing it (if it is not secreted). The selection of an appropriate host cell will depend upon various factors, such as desired expression levels, polypeptide modifications that are desirable or necessary for activity, such as glycosylation or phosphorylation, and ease of folding into a biologically active molecule.

A number of suitable host cells are known in the art and many are available from the American Type Culture Collection (ATCC), 10801 University Boulevard, Manassas, VA 20110-2209. Examples include, but are not limited to, mammalian cells, such as Chinese hamster ovary cells (CHO) (ATCC No. CCL61) CHO DHFR-cells (Urlaub et al., *Proc. Natl. Acad. Sci. USA*, 77:4216-4220 (1980)), human embryonic kidney (HEK) 293 or 293T cells (ATCC No. CRL1573), or 3T3 cells (ATCC No. CCL92). The selection of suitable mammalian host cells and methods for transformation, culture, amplification, screening and product production and purification are known in the art. Other suitable mammalian cell lines, are the monkey COS-1 (ATCC No. CRL1650) and COS-7 cell lines (ATCC No. CRL1651), and the CV-1 cell line (ATCC No. CCL70). Further exemplary mammalian host cells include primate cell lines and rodent cell lines, including transformed cell lines. Normal diploid cells, cell strains derived from *in vitro* culture of primary tissue, as well as primary explants, are also suitable. Candidate cells may be genotypically deficient in the selection gene, or may contain a dominantly acting selection gene. Other suitable mammalian cell lines include but are not limited to, mouse neuroblastoma N2A cells, HeLa, mouse L-929 cells, 3T3 lines derived from Swiss, Balb-c or NIH mice, BHK or HaK hamster cell lines, which are available from the ATCC. Each of these cell lines is known by and available to those skilled in the

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art of protein expression.

Similarly useful as host cells suitable for the present invention are bacterial cells. For example, the various strains of *E. coli* (e.g., HB101, (ATCC No. 33694) DH5 α , DH10, 5 and MC1061 (ATCC No. 53338)) are well-known as host cells in the field of biotechnology. Various strains of *B. subtilis*, *Pseudomonas spp.*, other *Bacillus spp.*, *Streptomyces spp.*, and the like may also be employed in this method.

Many strains of yeast cells known to those skilled in the 10 art are also available as host cells for the expression of the polypeptides of the present invention. Preferred yeast cells include, for example, *Saccharomyces cerevisiae* and *Pichia pastoris*.

Additionally, where desired, insect cell systems may be 15 utilized in the methods of the present invention. Such systems are described for example in Kitts et al., *Biotechniques*, 14:810-817 (1993); Lucklow, *Curr. Opin. Biotechnol.*, 4:564-572 (1993); and Lucklow et al. (*J. Virol.*, 67:4566-4579 (1993)). Preferred insect cells are Sf-9 and Hi5 20 (Invitrogen, Carlsbad, CA).

One may also use transgenic animals to express glycosylated IL-17 receptor like polypeptides. For example, one may use a transgenic milk-producing animal (a cow or goat, for example) and obtain the present glycosylated polypeptide 25 in the animal milk. One may also use plants to produce IL-17 receptor like polypeptides, however, in general, the glycosylation occurring in plants is different from that produced in mammalian cells, and may result in a glycosylated product which is not suitable for human therapeutic use.

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Polypeptide Production

Host cells comprising an IL-17 receptor like polypeptide expression vector may be cultured using standard media well known to the skilled artisan. The media will usually contain
5 all nutrients necessary for the growth and survival of the cells. Suitable media for culturing *E. coli* cells include, for example, Luria Broth (LB) and/or Terrific Broth (TB). Suitable media for culturing eukaryotic cells include Roswell Park Memorial Institute medium 1640 (RPMI 1640), Minimal
10 Essential Medium (MEM) and/or Dulbecco's Modified Eagle Medium (DMEM), all of which may be supplemented with serum and/or growth factors as indicated by the particular cell line being cultured. A suitable medium for insect cultures is Grace's medium supplemented with yeastolate, lactalbumin hydrolysate
15 and/or fetal calf serum, as necessary.

Typically, an antibiotic or other compound useful for selective growth of transformed cells is added as a supplement to the media. The compound to be used will be dictated by the selectable marker element present on the plasmid with which
20 the host cell was transformed. For example, where the selectable marker element is kanamycin resistance, the compound added to the culture medium will be kanamycin. Other compounds for selective growth include ampicillin, tetracycline, and neomycin.

25 The amount of an IL-17 receptor like polypeptide produced by a host cell can be evaluated using standard methods known in the art. Such methods include, without limitation, Western blot analysis, SDS-polyacrylamide gel electrophoresis, non-denaturing gel electrophoresis, high performance liquid
30 chromatography (HPLC) separation, immunoprecipitation, and/or activity assays such as DNA binding gel shift assays.

If an IL-17 receptor like polypeptide has been designed to be secreted from the host cells, the majority of polypeptide may be found in the cell culture medium. If

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however, the IL-17 receptor like polypeptide is not secreted from the host cells, it will be present in the cytoplasm and/or the nucleus (for eukaryotic host cells) or in the cytosol (for bacterial host cells).

5 For an IL-17 receptor like polypeptide situated in the host cell cytoplasm and/or the nucleus (for eukaryotic host cells) or in the cytosol (for bacterial host cells), intracellular material (including inclusion bodies for gram-negative bacteria) can be extracted from the host cell using
10 any standard technique known to the skilled artisan. For example, the host cells can be lysed to release the contents of the periplasm/cytoplasm by French press, homogenization, and/or sonication followed by centrifugation.

If an IL-17 receptor like polypeptide has formed
15 inclusion bodies in the cytosol, the inclusion bodies can often bind to the inner and/or outer cellular membranes and thus will be found primarily in the pellet material after centrifugation. The pellet material can then be treated at pH extremes or with a chaotropic agent such as a detergent,
20 guanidine, guanidine derivatives, urea, or urea derivatives in the presence of a reducing agent such as dithiothreitol at alkaline pH or tris carboxyethyl phosphine at acid pH to release, break apart, and solubilize the inclusion bodies. The IL-17 receptor like polypeptide in its now soluble form
25 can then be analyzed using gel electrophoresis, immunoprecipitation or the like. If it is desired to isolate the IL-17 receptor like polypeptide, isolation may be accomplished using standard methods such as those described herein and in Marston et al., *Meth. Enz.*, 182:264-275 (1990).

30 In some cases, an IL-17 receptor like polypeptide may not be biologically active upon isolation. Various methods for "refolding" or converting the polypeptide to its tertiary structure and generating disulfide linkages can be used to restore biological activity. Such methods include exposing

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the solubilized polypeptide to a pH usually above 7 and in the presence of a particular concentration of a chaotrope. The selection of chaotrope is very similar to the choices used for inclusion body solubilization, but usually the chaotrope is
5 used at a lower concentration and is not necessarily the same as chaotropes used for the solubilization. In most cases the refolding/oxidation solution will also contain a reducing agent or the reducing agent plus its oxidized form in a specific ratio to generate a particular redox potential
10 allowing for disulfide shuffling to occur in the formation of the protein's cysteine bridge(s). Some of the commonly used redox couples include cysteine/cystamine, glutathione (GSH)/dithiobis GSH, cupric chloride, dithiothreitol(DTT)/dithiane DTT, and 2-2mercaptoethanol(bME)/dithio-b(ME). A
15 cosolvent may be used to increase the efficiency of the refolding, and the more common reagents used for this purpose include glycerol, polyethylene glycol of various molecular weights, arginine and the like.

If inclusion bodies are not formed to a significant
20 degree upon expression of an IL-17 receptor like polypeptide, then the polypeptide will be found primarily in the supernatant after centrifugation of the cell homogenate. The polypeptide may be further isolated from the supernatant using methods such as those described herein.

The purification of an IL-17 receptor like polypeptide
25 from solution can be accomplished using a variety of techniques. If the polypeptide has been synthesized such that it contains a tag such as Hexahistidine (IL-17 receptor like polypeptide/hexaHis) or other small peptide such as FLAG
30 (Eastman Kodak Co., New Haven, CT) or myc (Invitrogen, Carlsbad, CA) at either its carboxyl or amino terminus, it may be purified in a one-step process by passing the solution through an affinity column where the column matrix has a high affinity for the tag.

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For example, polyhistidine binds with great affinity and specificity to nickel, thus an affinity column of nickel (such as the Qiagen® nickel columns) can be used for purification of IL-17 receptor like polypeptide/polyHis. See for example,
5 Ausubel et al., eds., *Current Protocols in Molecular Biology*, Section 10.11.8, John Wiley & Sons, New York (1993).

Additionally, the IL-17 receptor like polypeptide may be purified through the use of a monoclonal antibody which is capable of specifically recognizing and binding to the IL-17
10 receptor like polypeptide.

Suitable procedures for purification thus include, without limitation, affinity chromatography, immunoaffinity chromatography, ion exchange chromatography, molecular sieve chromatography, High Performance Liquid Chromatography (HPLC),
15 electrophoresis (including native gel electrophoresis) followed by gel elution, and preparative isoelectric focusing ("Isoprime" machine/technique, Hoefer Scientific, San Francisco, CA). In some cases, two or more purification techniques may be combined to achieve increased purity.

IL-17 receptor like polypeptides, including fragments, variants, and/or derivatives thereof may also be prepared by chemical synthesis methods (such as solid phase peptide synthesis) using techniques known in the art, such as those set forth by Merrifield et al., *J. Am. Chem. Soc.*, 85:2149
25 (1963), Houghten et al., *Proc Natl Acad. Sci. USA*, 82:5132 (1985), and Stewart and Young, *Solid Phase Peptide Synthesis*, Pierce Chemical Co., Rockford, IL (1984). Such polypeptides may be synthesized with or without a methionine on the amino terminus. Chemically synthesized IL-17 receptor like
30 polypeptides may be oxidized using methods set forth in these references to form disulfide bridges. Chemically synthesized IL-17 receptor like polypeptides are expected to have comparable biological activity to the corresponding IL-17

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receptor like polypeptides produced recombinantly or purified from natural sources, and thus may be used interchangeably with a recombinant or natural IL-17 receptor like polypeptide.

Another means of obtaining an IL-17 receptor like polypeptide is via purification from biological samples such as source tissues and/or fluids in which the IL-17 receptor like polypeptide is naturally found. Such purification can be conducted using methods for protein purification as described herein. The presence of the IL-17 receptor like polypeptide during purification may be monitored using, for example, an antibody prepared against recombinantly produced IL-17 receptor like polypeptide or peptide fragments thereof.

A number of additional methods for producing nucleic acids and polypeptides are known in the art, and can be used to produce polypeptides having specificity for IL-17 receptor like. See for example, Roberts et al., *Proc. Natl. Acad. Sci.*, 94:12297-12303 (1997), which describes the production of fusion proteins between an mRNA and its encoded peptide. See also Roberts, R., *Curr. Opin. Chem. Biol.*, 3:268-273 (1999). Additionally, U.S. patent No. 5,824,469 describes methods of obtaining oligonucleotides capable of carrying out a specific biological function. The procedure involves generating a heterogeneous pool of oligonucleotides, each having a 5' randomized sequence, a central preselected sequence, and a 3' randomized sequence. The resulting heterogeneous pool is introduced into a population of cells that do not exhibit the desired biological function. Subpopulations of the cells are then screened for those which exhibit a predetermined biological function. From that subpopulation, oligonucleotides capable of carrying out the desired biological function are isolated.

U.S. Patent Nos. 5,763,192, 5,814,476, 5,723,323, and 5,817,483 describe processes for producing peptides or polypeptides. This is done by producing stochastic genes or

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fragments thereof, and then introducing these genes into host cells which produce one or more proteins encoded by the stochastic genes. The host cells are then screened to identify those clones producing peptides or polypeptides
5 having the desired activity.

Another method for producing peptides or polypeptides is described in PCT/US98/20094 (WO99/15650) filed by Athersys, Inc. Known as "Random Activation of Gene Expression for Gene Discovery" (RAGE-GD), the process involves
10 the activation of endogenous gene expression or over-expression of a gene by *in situ* recombination methods. For example, expression of an endogenous gene is activated or increased by integrating a regulatory sequence into the target cell which is capable of activating expression of the gene by
15 non-homologous or illegitimate recombination. The target DNA is first subjected to radiation, and a genetic promoter inserted. The promoter eventually locates a break at the front of a gene, initiating transcription of the gene. This results in expression of the desired peptide or polypeptide.
20

It will be appreciated that these methods can also be used to create comprehensive IL-17 RECEPTOR like protein expression libraries, which can subsequently be used for high throughput phenotypic screening in a variety of assays, such as biochemical assays, cellular assays, and whole organism
25 assays (e.g., plant, mouse, etc.).

Chemical Derivatives

Chemically modified derivatives of the IL-17 receptor like polypeptides may be prepared by one skilled in the art, given the disclosures set forth hereinbelow. IL-17 receptor
30 like polypeptide derivatives are modified in a manner that is different, either in the type or location of the molecules naturally attached to the polypeptide. Derivatives may include molecules formed by the deletion of one or more naturally-attached chemical groups. The polypeptide

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comprising the amino acid sequence of SEQ ID NO: 2, or an IL-17 receptor like polypeptide variant may be modified by the covalent attachment of one or more polymers. For example, the polymer selected is typically water soluble so that the protein to which it is attached does not precipitate in an aqueous environment, such as a physiological environment. Included within the scope of suitable polymers is a mixture of polymers. Preferably, for therapeutic use of the end-product preparation, the polymer will be pharmaceutically acceptable.

The polymers each may be of any molecular weight and may be branched or unbranched. The polymers each typically have an average molecular weight of between about 2kDa to about 100kDa (the term "about" indicating that in preparations of a water soluble polymer, some molecules will weigh more, some less, than the stated molecular weight). The average molecular weight of each polymer preferably is between about 5kDa and about 50kDa, more preferably between about 12kDa and about 40kDa and most preferably between about 20kDa and about 35kDa.

Suitable water soluble polymers or mixtures thereof include, but are not limited to, N-linked or O-linked carbohydrates, sugars, phosphates, polyethylene glycol (PEG) (including the forms of PEG that have been used to derivatize proteins, including mono-(C₁-C₁₀) alkoxy- or aryloxy-polyethylene glycol), monomethoxy-polyethylene glycol, dextran (such as low molecular weight dextran, of, for example about 6 kD), cellulose, or other carbohydrate based polymers, poly-(N-vinyl pyrrolidone) polyethylene glycol, propylene glycol homopolymers, a polypropylene oxide/ethylene oxide co-polymer, polyoxyethylated polyols (e.g., glycerol) and polyvinyl alcohol. Also encompassed by the present invention are bifunctional crosslinking molecules which may be used to prepare covalently attached multimers of the polypeptide

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comprising the amino acid sequence of SEQ ID NO: 2 or an IL-17 receptor like polypeptide variant.

In general, chemical derivatization may be performed under any suitable condition used to react a protein with an activated polymer molecule. Methods for preparing chemical derivatives of polypeptides will generally comprise the steps of (a) reacting the polypeptide with the activated polymer molecule (such as a reactive ester or aldehyde derivative of the polymer molecule) under conditions whereby the polypeptide comprising the amino acid sequence of SEQ ID NO: 2, or an IL-17 receptor like polypeptide variant becomes attached to one or more polymer molecules, and (b) obtaining the reaction product(s). The optimal reaction conditions will be determined based on known parameters and the desired result.

For example, the larger the ratio of polymer molecules:protein, the greater the percentage of attached polymer molecule. In one embodiment, the IL-17 receptor like polypeptide derivative may have a single polymer molecule moiety at the amino terminus. See, for example, U.S. Patent No. 5,234,784.

The pegylation of the polypeptide specifically may be carried out by any of the pegylation reactions known in the art, as described for example in the following references: Francis et al., *Focus on Growth Factors*, 3:4-10 (1992); EP 0154316; EP 0401384 and U.S. Patent No. 4,179,337. For example, pegylation may be carried out via an acylation reaction or an alkylation reaction with a reactive polyethylene glycol molecule (or an analogous reactive water-soluble polymer) as described herein. For the acylation reactions, the polymer(s) selected should have a single reactive ester group. For reductive alkylation, the polymer(s) selected should have a single reactive aldehyde group. A reactive aldehyde is, for example, polyethylene glycol propionaldehyde, which is water stable, or mono-C₁-C₁₀

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alkoxy or aryloxy derivatives thereof (see U.S. Patent No. 5,252,714).

In another embodiment, IL-17 receptor like polypeptides may be chemically coupled to biotin, and the biotin/IL-17 receptor like polypeptide molecules which are conjugated are then allowed to bind to avidin, resulting in tetravalent avidin/biotin/IL-17 receptor like polypeptide molecules. IL-17 receptor like polypeptides may also be covalently coupled to dinitrophenol (DNP) or trinitrophenol (TNP) and the resulting conjugates precipitated with anti-DNP or anti-TNP-IgM to form decameric conjugates with a valency of 10.

Generally, conditions which may be alleviated or modulated by the administration of the present IL-17 receptor like polypeptide derivatives include those described herein for IL-17 receptor like polypeptides. However, the IL-17 receptor like polypeptide derivatives disclosed herein may have additional activities, enhanced or reduced biological activity, or other characteristics, such as increased or decreased half-life, as compared to the non-derivatized molecules.

The present invention further includes non-human animals in which the promoter for one or more of the IL-17 receptor like polypeptides of the present invention is either activated or inactivated (e.g., by using homologous recombination methods) to alter the level of expression of one or more of the native IL-17 receptor like polypeptides.

These non-human animals may be used for drug candidate screening. In such screening, the impact of a drug candidate on the animal may be measured. For example, drug candidates may decrease or increase the expression of the IL-17 receptor like gene. In certain embodiments, the amount of IL-17 receptor like polypeptide, that is produced may be measured

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after the exposure of the animal to the drug candidate. Additionally, in certain embodiments, one may detect the actual impact of the drug candidate on the animal. For example, the overexpression of a particular gene may result 5 in, or be associated with, a disease or pathological condition. In such cases, one may test a drug candidate's ability to decrease expression of the gene or its ability to prevent or inhibit a pathological condition. In other examples, the production of a particular metabolic product 10 such as a fragment of a polypeptide, may result in, or be associated with, a disease or pathological condition. In such cases, one may test a drug candidate's ability to decrease the production of such a metabolic product or its ability to prevent or inhibit a pathological condition.

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Microarray

It will be appreciated that DNA microarray technology can be utilized in accordance with the present invention. DNA microarrays are miniature, high density arrays of nucleic acids positioned on a solid support, such as glass. Each cell or element within the array has numerous copies of a single species of DNA which acts as a target for hybridization for its cognate mRNA. In expression profiling using DNA microarray technology, mRNA is first extracted from a cell or 20 tissue sample and then converted enzymatically to fluorescently labeled cDNA. This material is hybridized to the microarray and unbound cDNA is removed by washing. The expression of discrete genes represented on the array is then visualized by quantitating the amount of labeled cDNA which is 25 specifically bound to each target DNA. In this way, the expression of thousands of genes can be quantitated in a high throughput, parallel manner from a single sample of biological material.

This high throughput expression profiling has a broad 35 range of applications with respect to the IL-17 receptor like

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molecules of the invention, including, but not limited to: the identification and validation of IL-17 receptor like disease-related genes as targets for therapeutics; molecular toxicology of IL-17 receptor like molecules and inhibitors thereof; stratification of populations and generation of surrogate markers for clinical trials; and enhancing IL-17 receptor like-related small molecule drug discovery by aiding in the identification of selective compounds in high throughput screens (HTS).

10 **Selective Binding Agents**

As used herein, the term "selective binding agent" refers to a molecule which has specificity for one or more IL-17 receptor like polypeptides. Suitable selective binding agents include, but are not limited to, antibodies and derivatives thereof, polypeptides, and small molecules. Suitable selective binding agents may be prepared using methods known in the art. An exemplary IL-17 receptor like polypeptide selective binding agent of the present invention is capable of binding a certain portion of the IL-17 receptor like polypeptide thereby inhibiting the binding of the polypeptide to the IL-17 receptor like polypeptide receptor(s).

Selective binding agents such as antibodies and antibody fragments that bind IL-17 receptor like polypeptides are within the scope of the present invention. The antibodies may be polyclonal including monospecific polyclonal, monoclonal (MAbs), recombinant, chimeric, humanized such as CDR-grafted, human, single chain, and/or bispecific, as well as fragments, variants or derivatives thereof. Antibody fragments include those portions of the antibody which bind to an epitope on the IL-17 receptor like polypeptide. Examples of such fragments include Fab and F(ab') fragments generated by enzymatic cleavage of full-length antibodies. Other binding fragments include those generated by recombinant DNA techniques, such as

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the expression of recombinant plasmids containing nucleic acid sequences encoding antibody variable regions.

Polyclonal antibodies directed toward an IL-17 receptor like polypeptide generally are produced in animals (e.g., 5 rabbits or mice) by means of multiple subcutaneous or intraperitoneal injections of IL-17 receptor like polypeptide and an adjuvant. It may be useful to conjugate an IL-17 receptor like polypeptide, including a variant, fragment, or derivative, to a carrier protein that is immunogenic in the 10 species to be immunized, such as keyhole limpet hemocyanin, serum, albumin, bovine thyroglobulin, or soybean trypsin inhibitor. Also, aggregating agents such as alum are used to enhance the immune response. After immunization, the animals are bled and the serum is assayed for anti-IL-17 receptor like 15 polypeptide antibody titer.

Monoclonal antibodies directed toward an IL-17 receptor like polypeptide are produced using any method which provides for the production of antibody molecules by continuous cell lines in culture. Examples of suitable methods for preparing 20 monoclonal antibodies include the hybridoma methods of Kohler et al., *Nature*, 256:495-497 (1975) and the human B-cell hybridoma method, Kozbor, *J. Immunol.*, 133:3001 (1984); Brodeur et al., *Monoclonal Antibody Production Techniques and Applications*, pp. 51-63 (Marcel Dekker, Inc., New York, 1987). 25 Also provided by the invention are hybridoma cell lines which produce monoclonal antibodies reactive with IL-17 receptor like polypeptides.

Monoclonal antibodies of the invention may be modified for use as therapeutics. One embodiment is a "chimeric" 30 antibody in which a portion of the heavy and/or light chain is identical with or homologous to a corresponding sequence in antibodies derived from a particular species or belonging to a particular antibody class or subclass, while the remainder of the chain(s) is identical with or homologous to a

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corresponding sequence in antibodies derived from another species or belonging to another antibody class or subclass. Also included are fragments of such antibodies, so long as they exhibit the desired biological activity. See, U.S. 5 Patent No. 4,816,567; Morrison et al., *Proc. Natl. Acad. Sci.*, 81:6851-6855 (1985).

In another embodiment, a monoclonal antibody of the invention is a "humanized" antibody. Methods for humanizing non-human antibodies are well known in the art. See U.S. 10 Patent Nos. 5,585,089, and 5,693,762. Generally, a humanized antibody has one or more amino acid residues introduced into it from a source which is non-human. Humanization can be performed, for example, using methods described in the art (Jones et al., *Nature* 321:522-525 (1986); Riechmann et al., 15 *Nature*, 332:323-327 (1988); Verhoeyen et al., *Science* 239:1534-1536 (1988)), by substituting at least a portion of a rodent complementarity-determining region (CDR) for the corresponding regions of a human antibody.

Also encompassed by the invention are human antibodies 20 which bind IL-17 receptor like polypeptides. Using transgenic animals (e.g., mice) that are capable of producing a repertoire of human antibodies in the absence of endogenous immunoglobulin production such antibodies are produced by immunization with an IL-17 receptor like antigen (i.e., having 25 at least 6 contiguous amino acids), optionally conjugated to a carrier. See, for example, Jakobovits et al., *Proc. Natl. Acad. Sci.*, 90:2551-2555 (1993); Jakobovits et al., *Nature* 362:255-258 (1993); Brugermann et al., *Year in Immuno.*, 7:33 (1993). In one method, such transgenic animals are produced 30 by incapacitating the endogenous loci encoding the heavy and light immunoglobulin chains therein, and inserting loci encoding human heavy and light chain proteins into the genome thereof. Partially modified animals, that is those having less than the full complement of modifications, are then

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cross-bred to obtain an animal having all of the desired immune system modifications. When administered an immunogen, these transgenic animals produce antibodies with human variable regions, including human (rather than e.g., murine) 5 amino acid sequences, including variable regions, including human which are immunospecific for these antigens. See PCT application nos. PCT/US96/05928 and PCT/US93/06926. Additional methods are described in U.S. Patent No. 5,545,807, PCT application nos. PCT/US91/245, PCT/GB89/01207, and in EP 10 546073B1 and EP 546073A1. Human antibodies may also be produced by the expression of recombinant DNA in host cells or by expression in hybridoma cells as described herein.

In an alternative embodiment, human antibodies can be produced from phage-display libraries (Hoogenboom et al., J. 15 *Mol. Biol.* 227:381 (1991); Marks et al., *J. Mol. Biol.* 222:581 (1991). These processes mimic immune selection through the display of antibody repertoires on the surface of filamentous bacteriophage, and subsequent selection of phage by their binding to an antigen of choice. One such technique is 20 described in PCT Application no. PCT/US98/17364, which describes the isolation of high affinity and functional agonistic antibodies for MPL- and msk- receptors using such an approach.

Chimeric, CDR grafted, and humanized antibodies are 25 typically produced by recombinant methods. Nucleic acids encoding the antibodies are introduced into host cells and expressed using materials and procedures described herein. In a preferred embodiment, the antibodies are produced in mammalian host cells, such as CHO cells. Monoclonal (e.g., 30 human) antibodies may be produced by the expression of recombinant DNA in host cells or by expression in hybridoma cells as described herein.

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- The anti-IL-17 receptor like antibodies of the invention may be employed in any known assay method, such as competitive binding assays, direct and indirect sandwich assays, and immunoprecipitation assays (Sola, Monoclonal Antibodies: A Manual of Techniques, pp. 147-158 (CRC Press, Inc., 1987)) for the detection and quantitation of IL-17 receptor like polypeptides. The antibodies will bind IL-17 receptor like polypeptides with an affinity which is appropriate for the assay method being employed.
- For diagnostic applications, in certain embodiments, anti-IL-17 receptor like antibodies may be labeled with a detectable moiety. The detectable moiety can be any one which is capable of producing, either directly or indirectly, a detectable signal. For example, the detectable moiety may be a radioisotope, such as ^3H , ^{14}C , ^{32}P , ^{35}S , or ^{125}I , a fluorescent or chemiluminescent compound, such as fluorescein isothiocyanate, rhodamine, or luciferin; or an enzyme, such as alkaline phosphatase, β -galactosidase, or horseradish peroxidase (Bayer et al., *Meth. Enz.*, 184:138-163 (1990)).
- Competitive binding assays rely on the ability of a labeled standard (e.g., an IL-17 receptor like polypeptide, or an immunologically reactive portion thereof) to compete with the test sample analyte (an IL-17 receptor like polypeptide) for binding with a limited amount of anti IL-17 receptor like antibody. The amount of an IL-17 receptor like polypeptide in the test sample is inversely proportional to the amount of standard that becomes bound to the antibodies. To facilitate determining the amount of standard that becomes bound, the antibodies typically are insolubilized before or after the competition, so that the standard and analyte that are bound to the antibodies may conveniently be separated from the standard and analyte which remain unbound.

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Sandwich assays typically involve the use of two antibodies, each capable of binding to a different immunogenic portion, or epitope, of the protein to be detected and/or quantitated. In a sandwich assay, the test sample analyte is 5 typically bound by a first antibody which is immobilized on a solid support, and thereafter a second antibody binds to the analyte, thus forming an insoluble three part complex. See, e.g., U.S. Patent No. 4,376,110. The second antibody may itself be labeled with a detectable moiety (direct sandwich 10 assays) or may be measured using an anti-immunoglobulin antibody that is labeled with a detectable moiety (indirect sandwich assays). For example, one type of sandwich assay is an enzyme-linked immunosorbent assay (ELISA), in which case the detectable moiety is an enzyme.

15 The selective binding agents, including anti-IL-17 receptor like antibodies, also are useful for *in vivo* imaging. An antibody labeled with a detectable moiety may be administered to an animal, preferably into the bloodstream, and the presence and location of the labeled antibody in the 20 host is assayed. The antibody may be labeled with any moiety that is detectable in an animal, whether by nuclear magnetic resonance, radiology, or other detection means known in the art.

The invention also relates to a kit comprising IL-17 25 receptor like selective binding agents (such as antibodies) and other reagents useful for detecting IL-17 receptor like polypeptide levels in biological samples. Such reagents may include a secondary activity, a detectable label, blocking serum, positive and negative control samples, and detection 30 reagents

Selective binding agents of the invention, including antibodies, may be used as therapeutics. These therapeutic agents are generally agonists or antagonists, in that they

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either enhance or reduce, respectively, at least one of the biological activities of an IL-17 receptor like polypeptide. In one embodiment, antagonist antibodies of the invention are antibodies or binding fragments thereof which are capable of 5 specifically binding to an IL-17 receptor like polypeptide and which are capable of inhibiting or eliminating the functional activity of an IL-17 receptor like polypeptide *in vivo* or *in vitro*. In preferred embodiments, the selective binding agent, e.g., an antagonist antibody, will inhibit the functional 10 activity of an IL-17 receptor like polypeptide by at least about 50%, and preferably by at least about 80%. In another embodiment, the selective binding agent may be an anti-IL-17 receptor like polypeptide antibody that is capable of interacting with an IL-17 receptor like binding partner (a 15 ligand or receptor) thereby inhibiting or eliminating IL-17 receptor like activity *in vitro* or *in vivo*. Selective binding agents, including agonist and antagonist anti-IL-17 receptor like antibodies, are identified by screening assays which are well known in the art.

20 The invention also relates to a kit comprising IL-17 receptor like selective binding agents (such as antibodies) and other reagents useful for detecting IL-17 receptor like polypeptide levels in biological samples. Such reagents may include, a detectable label, blocking serum, positive and 25 negative control samples, and detection reagents.

IL-17 receptor like polypeptides can be used to clone IL-17 receptor like ligand(s) using an "expression cloning" strategy. Radiolabeled (125-Iodine) IL-17 receptor like polypeptide or "affinity/activity-tagged" IL-17 receptor like 30 polypeptide (such as an Fc fusion or an alkaline phosphatase fusion) can be used in binding assays to identify a cell type or cell line or tissue that expresses IL-17 receptor like ligand(s). RNA isolated from such cells or tissues can then be converted to cDNA, cloned into a mammalian expression

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vector, and transfected into mammalian cells (for example, COS, or 293) to create an expression library. Radiolabeled or tagged IL-17 receptor like polypeptide can then be used as an affinity reagent to identify and isolate the subset of cells 5 in this library expressing IL-17 receptor like ligand(s). DNA is then isolated from these cells and transfected into mammalian cells to create a secondary expression library in which the fraction of cells expressing IL-17 receptor like ligand(s) would be many-fold higher than in the original 10 library. This enrichment process can be repeated iteratively until a single recombinant clone containing an IL-17 receptor like ligand is isolated. Isolation of IL-17 receptor like ligand(s) is useful for identifying or developing novel agonists and antagonists of the IL-17 receptor like signaling 15 pathway. Such agonists and antagonists include IL-17 receptor like ligand(s), anti-IL-17 receptor like ligand antibodies, small molecules or antisense oligonucleotides.

20 Assaying for Other Modulators of IL-17 Receptor Like Polypeptide Activity

In some situations, it may be desirable to identify molecules that are modulators, i.e., agonists or antagonists, of the activity of IL-17 receptor like polypeptide. Natural or synthetic molecules that modulate IL-17 receptor like 25 polypeptide may be identified using one or more screening assays, such as those described herein. Such molecules may be administered either in an *ex vivo* manner, or in an *in vivo* manner by injection, or by oral delivery, implantation device, or the like.

30 "Test molecule(s)" refers to the molecule(s) that is/are under evaluation for the ability to modulate (i.e., increase or decrease) the activity of an IL-17 receptor like polypeptide. Most commonly, a test molecule will interact directly with an IL-17 receptor like polypeptide. However, it

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is also contemplated that a test molecule may also modulate IL-17 receptor like polypeptide activity indirectly, such as by affecting IL-17 receptor like gene expression, or by binding to an IL-17 receptor like binding partner (e.g., 5 receptor or ligand). In one embodiment, a test molecule will bind to an IL-17 receptor like polypeptide with an affinity constant of at least about 10^{-6} M, preferably about 10^{-8} M, more preferably about 10^{-9} M, and even more preferably about 10^{-10} M.

10 Methods for identifying compounds which interact with IL-17 receptor like polypeptides are encompassed by the present invention. In certain embodiments, an IL-17 receptor like polypeptide is incubated with a test molecule under conditions which permit the interaction of the test molecule with an IL-15 17 receptor like polypeptide, and the extent of the interaction can be measured. The test molecule(s) can be screened in a substantially purified form or in a crude mixture. The test molecules can be nucleic acids, proteins, peptides, carbohydrates, lipids, organic and inorganic 20 compounds.

In certain embodiments, an IL-17 receptor like polypeptide agonist or antagonist may be a protein, peptide, carbohydrate, lipid, or small molecular weight molecule which interacts with IL-17 receptor like polypeptide, or ligand 25 thereof, to regulate its activity. Molecules which regulate IL-17 receptor like polypeptide expression include nucleic acids which are complementary to nucleic acids encoding an IL-17 receptor like polypeptide, or are complementary to nucleic acids sequences which direct or control the expression of IL-30 17 receptor like polypeptide, and which act as anti-sense regulators of expression.

Once a set of test molecules has been identified as interacting with an IL-17 receptor like polypeptide, the

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molecules may be further evaluated for their ability to increase or decrease IL-17 receptor like polypeptide activity. The measurement of the interaction of test molecules with IL-17 receptor like polypeptides may be carried out in several 5 formats, including cell-based binding assays, membrane binding assays, solution-phase assays and immunoassays. In general, test molecules are incubated with an IL-17 receptor like polypeptide for a specified period of time, and IL-17 receptor like polypeptide activity is determined by one or more assays 10 for measuring biological activity.

The interaction of test molecules with IL-17 receptor like polypeptides may also be assayed directly using polyclonal or monoclonal antibodies in an immunoassay. Alternatively, modified forms of IL-17 receptor like 15 polypeptides containing epitope tags as described herein may be used in immunoassays.

In certain embodiments, a IL-17 receptor like polypeptide agonist or antagonist may be a protein, peptide, carbohydrate, lipid, or small molecular weight molecule which interacts with 20 IL-17 receptor like polypeptide to regulate its activity. Potential protein antagonists of IL-17 receptor like polypeptide include antibodies which interact with active regions of the polypeptide and inhibit or eliminate at least one activity of IL-17 receptor like molecules. Molecules 25 which regulate IL-17 receptor like polypeptide expression include nucleic acids which are complementary to nucleic acids encoding a IL-17 receptor like polypeptide, or are complementary to nucleic acids sequences which direct or control the expression of IL-17 receptor like polypeptide, and 30 which act as anti-sense regulators of expression.

In the event that IL-17 receptor like polypeptides display biological activity through an interaction with a binding partner (e.g., selective binding agent or a ligand), a

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variety of *in vitro* assays may be used to measure the binding of an IL-17 receptor like polypeptide to the corresponding binding partner (such as a selective binding agent or ligand). These assays may be used to screen test molecules for their ability to increase or decrease the rate and/or the extent of binding of an IL-17 receptor like polypeptide to its binding partner. In one assay, an IL-17 receptor like polypeptide is immobilized in the wells of a microtiter plate. Radiolabeled IL-17 receptor like binding partner (for example, iodinated IL-17 receptor like binding partner) and the test molecule(s) can then be added either one at a time (in either order) or simultaneously to the wells. After incubation, the wells can be washed and counted, using a scintillation counter, for radioactivity to determine the extent to which the binding partner bound to IL-17 receptor like polypeptide. Typically, the molecules will be tested over a range of concentrations, and a series of control wells lacking one or more elements of the test assays can be used for accuracy in the evaluation of the results. An alternative to this method involves reversing the "positions" of the proteins, i.e., immobilizing IL-17 receptor like binding partner to the microtiter plate wells, incubating with the test molecule and radiolabeled IL-17 receptor like polypeptide, and determining the extent of IL-17 receptor like polypeptide binding. See, for example, chapter 18, *Current Protocols in Molecular Biology*, Ausubel et al., eds., John Wiley & Sons, New York, NY (1995).

As an alternative to radiolabelling, an IL-17 receptor like polypeptide or its binding partner may be conjugated to biotin and the presence of biotinylated protein can then be detected using streptavidin linked to an enzyme, such as horseradish peroxidase (HRP) or alkaline phosphatase (AP), that can be detected colorometrically, or by fluorescent tagging of streptavidin. An antibody directed to an IL-17 receptor like polypeptide or to an IL-17 receptor like binding partner and conjugated to biotin may also be used and can be

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detected after incubation with enzyme-linked streptavidin linked to AP or HRP.

An IL-17 receptor like polypeptide or an IL-17 receptor like binding partner can also be immobilized by attachment to 5 agarose beads, acrylic beads or other types of such inert solid phase substrates. The substrate-protein complex can be placed in a solution containing the complementary protein and the test compound. After incubation, the beads can be precipitated by centrifugation, and the amount of binding 10 between an IL-17 receptor like polypeptide and its binding partner can be assessed using the methods described herein. Alternatively, the substrate-protein complex can be immobilized in a column, and the test molecule and 15 complementary protein are passed through the column. The formation of a complex between an IL-17 receptor like polypeptide and its binding partner can then be assessed using any of the techniques set forth herein, i.e., radiolabelling, antibody binding, or the like.

Another *in vitro* assay that is useful for identifying a 20 test molecule which increases or decreases the formation of a complex between an IL-17 receptor like binding protein and an IL-17 receptor like binding partner is a surface plasmon resonance detector system such as the BIACore assay system (Pharmacia, Piscataway, NJ). The BIACore system may be 25 carried out using the manufacturer's protocol. This assay essentially involves the covalent binding of either IL-17 receptor like polypeptide or an IL-17 receptor like binding partner to a dextran-coated sensor chip which is located in a detector. The test compound and the other complementary 30 protein can then be injected, either simultaneously or sequentially, into the chamber containing the sensor chip. The amount of complementary protein that binds can be assessed based on the change in molecular mass which is physically associated with the dextran-coated side of the sensor chip;

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the change in molecular mass can be measured by the detector system.

In some cases, it may be desirable to evaluate two or more test compounds together for their ability to increase or

5 decrease the formation of a complex between an IL-17 receptor like polypeptide and an IL-17 receptor like binding partner.

In these cases, the assays set forth herein can be readily modified by adding such additional test compound(s) either simultaneous with, or subsequent to, the first test compound.

10 The remainder of the steps in the assay are as set forth herein.

In vitro assays such as those described herein may be used advantageously to screen large numbers of compounds for effects on complex formation by IL-17 receptor like

15 polypeptide and IL-17 receptor like binding partner. The assays may be automated to screen compounds generated in phage display, synthetic peptide, and chemical synthesis libraries.

Compounds which increase or decrease the formation of a complex between an IL-17 receptor like polypeptide and an IL-

20 17 receptor like binding partner may also be screened in cell culture using cells and cell lines expressing either IL-17 receptor like polypeptide or IL-17 receptor like binding partner.

Cells and cell lines may be obtained from any mammal, but preferably will be from human or other primate,

25 canine, or rodent sources. The binding of an IL-17 receptor like polypeptide to cells expressing IL-17 receptor like binding partner at the surface is evaluated in the presence or absence of test molecules, and the extent of binding may be determined by, for example, flow cytometry using a

30 biotinylated antibody to an IL-17 receptor like binding partner. Cell culture assays can be used advantageously to further evaluate compounds that score positive in protein

binding assays described herein.

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Cell cultures can also be used to screen the impact of a drug candidate. For example, drug candidates may decrease or increase the expression of the IL-17 receptor like gene. In certain embodiments, the amount of IL-17 receptor like polypeptide or a fragment that is produced may be measured after exposure of the cell culture to the drug candidate. In certain embodiments, one may detect the actual impact of the drug candidate on the cell culture. For example, the overexpression of a particular gene may have a particular impact on the cell culture. In such cases, one may test a drug candidate's ability to increase or decrease the expression of the gene or its ability to prevent or inhibit a particular impact on the cell culture. In other examples, the production of a particular metabolic product such as a fragment of a polypeptide, may result in, or be associated with, a disease or pathological condition. In such cases, one may test a drug candidate's ability to decrease the production of such a metabolic product in a cell culture.

A yeast two hybrid system (Chien et al., Proc. Natl. Acad. Sci. USA, 88: 9578-9583, 1991) can be used to identify novel polypeptides that bind to a yeast-two hybrid bait construct can be generated in a vector(such as the pAS2-1 form Clontech) which encodes a yeast-two hybrid domain fused to the IL-17 receptor like polynucleotide.. This bait construct may be used to screen human cDNA libraries wherein the cDNA library sequences are fused to GAL4 activation domains. Positive interactions will result in the activation of a reporter gene such as β -gal. Positive clones emerging from the screening may be characterized further to identify interacting proteins.

Internalizing Proteins

The tat protein sequence (from HIV) can be used to internalize proteins into a cell. See e.g., Falwell et al.,

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Proc. Natl. Acad. Sci., 91:664-668 (1994). For example, an 11 amino acid sequence (YGRKKRQRR; SEQ ID NO: 16) of the HIV tat protein (termed the "protein transduction domain", or TAT PDT) has been described as mediating delivery across the cytoplasmic membrane and the nuclear membrane of a cell. See Schwarze et al., *Science*, 285:1569-1572 (1999); and Nagahara et al., *Nature Medicine*, 4:1449-1452 (1998). In these procedures, FITC-constructs (FITC-GGGGYGRKKRQRR; SEQ ID NO: 17) are prepared which bind to cells as observed by fluorescence-activated cell sorting (FACS) analysis, and these constructs penetrate tissues after i.p. administration. Next, tat-bgal fusion proteins are constructed. Cells treated with this construct demonstrated β -gal activity. Following injection, a number of tissues, including liver, kidney, lung, heart, and brain tissue have been found to demonstrate expression using these procedures. It is believed that these constructions underwent some degree of unfolding in order to enter the cell; as such, refolding may be required after entering the cell.

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It will thus be appreciated that the tat protein sequence may be used to internalize a desired protein or polypeptide into a cell. For example, using the tat protein sequence, an IL-17 receptor like antagonist (such as an anti-IL-17 receptor like selective binding agent, small molecule, soluble receptor, or antisense oligonucleotide) can be administered intracellularly to inhibit the activity of an IL-17 receptor like molecule. As used herein, the term "IL-17 receptor like molecule" refers to both IL-17 receptor like nucleic acid molecules and IL-17 receptor like polypeptides as defined herein. Where desired, the IL-17 receptor like protein itself may also be internally administered to a cell using these procedures. See also, Strauss, E., "Introducing Proteins Into the Body's Cells", *Science*, 285:1466-1467 (1999).

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Cell Source Identification Using IL-17 Receptor Like Polypeptides

In accordance with certain embodiments of the invention, it may be useful to be able to determine the source of a 5 certain cell type associated with an IL-17 receptor like polypeptide. For example, it may be useful to determine the origin of a disease or pathological condition as an aid in selecting an appropriate therapy.

Therapeutic Uses

10 A non-exclusive list of acute and chronic diseases which can be treated, diagnosed, ameliorated, or prevented with the IL-17 receptor like nucleic acids, polypeptides, and agonists and antagonists of the invention include:

- The diagnosis and/or treatment of diseases involving immune system dysfunction. Examples of such diseases include, but are not limited to, rheumatoid arthritis, psoriatic arthritis, inflammatory arthritis, osteoarthritis, inflammatory joint disease, autoimmune disease including autoimmune vasculitis, multiple sclerosis, lupus, diabetes (e.g., insulin diabetes), inflammatory bowel disease, transplant rejection, graft vs. host disease, and inflammatory conditions resulting from strain, sprain, cartilage damage, trauma, orthopedic surgery, infection or other disease processes. Other 20 diseases influenced by the dysfunction of the immune system are encompassed within the scope of the invention, including but not limited to, allergies. The IL-17 receptor like nucleic acids, polypeptides, and agonists and antagonists of the invention can also be used to 25 inhibit T cell proliferation, to inhibit T cell activation, and/or to inhibit B cell proliferation and/or immunoglobulin secretion.

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- The diagnosis and/or treatment of diseases involving infection. Examples of such diseases include, but are not limited to, leprosy, viral infections such as hepatitis or HIV, bacterial infection such as *clostridium* associated illnesses, including *clostridium*-associated diarrhea, pulmonary tuberculosis, acute febrile illness from bacteria such as or virus, fever, acute phase response of the liver, septicemia, septic shock. Other diseases involving infection are encompassed within the scope of the invention.
- The diagnosis and/or treatment of diseases involving weight disorders. Examples of such diseases include, but are not limited to obesity, anorexia, cachexia, including AIDS-induced cachexia, myopathies (e.g., muscle protein metabolism, such as in sepsis), and hypoglycemia. Other diseases involving weight disorders are encompassed within the scope of the invention.
- The diagnosis and/or treatment of diseases involving neuronal dysfunction. Examples of such diseases include, but are not limited to Alzheimer's, Parkinson's disease, neurotoxicity (e.g., as induced by HIV), ALS, brain injury, stress, depression, nociception and other pain (including cancer-related pain), hyperalgesia, epilepsy, learning impairment and memory disorders, sleep disturbance, and peripheral and central neuropathies. Other neurological disorders are encompassed within the scope of the invention.
- The diagnosis and/or treatment of diseases involving the lung. Examples of such diseases include, but are not limited to, acute or chronic lung injury including interstitial lung disease, acute respiratory disease syndrome, pulmonary hypertension, emphysema, cystic fibrosis, pulmonary fibrosis, and asthma. Other diseases of the lung are encompassed within the scope of the

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invention.

- The diagnosis and/or treatment of diseases involving the skin. Examples of such diseases include, but are not limited to, psoriasis, eczema, and wound healing. Other diseases of the skin are encompassed within the scope of the invention.
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- The diagnosis and/or treatment of diseases involving the kidney. Examples of such diseases include, but are not limited to, acute and chronic glomerulonephritis. Other diseases of the kidney are encompassed within the scope
10 of the invention.
- The diagnosis and/or treatment of diseases involving the bone. Examples of such diseases include, but are not limited to, osteoporosis, osteopetrosis, osteogenesis imperfecta, Paget's disease, periodontal disease, temporal mandibular joint disease, and hypercalcemia. Other diseases of the bone are encompassed within the
15 scope of the invention.
- 20 • The diagnosis and/or treatment of diseases involving the vascular system. Examples of such diseases include, but are not limited to hemorrhage or stroke, hemorrhagic shock, ischemia, including cardiac ischemia and cerebral ischemia (e.g., brain injury as a result of trauma, epilepsy, hemorrhage or stroke, each of which may lead to neurodegeneration), atherosclerosis, congestive heart failure; restenosis, reperfusion injury, and angiogenesis. Other diseases of the vascular system are encompassed within the scope of the invention.
25
- 30 • The diagnosis and/or treatment of tumor cells. Examples of such diseases include, but are not limited to, lymphomas, bone sarcoma, chronic and acute myelogenous leukemia (AML and CML) and other leukemias, multiple

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myeloma, lung, breast cancer, tumor metastasis, and side effects from radiation therapy. Other diseases involving tumor cells are encompassed within the scope of the invention.

- 5 • The diagnosis and/or treatment of reproductive disorders. Examples of such diseases include, but are not limited to, infertility, miscarriage, pre-term labor and delivery, and endometriosis. Other diseases involving the reproductive system are encompassed within the scope of .
10 the invention.
- The diagnosis and/or treatment of eye disorders. Examples of such diseases include, but are not limited to, inflammatory eye disease, as may be associated with, for example, corneal transplant; retinal degeneration,
15 blindness, macular degeneration, glaucoma, uveitis, and retinal neuropathy. Other diseases of the eye are encompassed within the scope of the invention.

20 Other diseases which are treatable using agents within the scope of the invention include acute pancreatitis, chronic fatigue syndrome, fibromyalgia, and Kawasaki's disease (MLNS).

Other diseases associated with undesirable levels of one or more of IL-1, IL-1ra, the ligand of the present IL-17
25 receptor like polypeptide, and/or the present IL-17 receptor like polypeptide itself are encompassed within the scope of .
the invention. Undesirable levels include excessive and/or sub-normal levels of IL-1, IL-1ra, the ligand of the present
30 IL-17 receptor like polypeptide, and/or the IL-17 receptor like polypeptides described herein.

IL-1 inhibitors include any protein capable of specifically preventing activation of cellular receptors to

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IL-1, which may result from any number of mechanisms. Such mechanisms include downregulating IL-1 production, binding free IL-1, interfering with IL-1 binding to its receptor, interfering with formation of the IL-1 receptor complex (i.e.,
5 association of IL-1 receptor with IL-1 receptor accessory protein), or interfering with modulation of IL-1 signaling after binding to its receptor. Classes of interleukin-1 inhibitors include:

- Interleukin-1 receptor antagonists such as IL-1ra, as
10 described herein;
- Anti-IL-1 receptor monoclonal antibodies (e.g., EP 623674);
- IL-1 binding proteins such as soluble IL-1 receptors (e.g., U.S. Pat. No. 5,492,888, U.S. Pat. No. 5,488,032,
15 and U.S. Pat. No. 5,464,937, U.S. Pat. No. 5,319,071, and U.S. Pat. No. 5,180,812);
- Anti-IL-1 monoclonal antibodies (e.g., WO 9501997, WO 9402627, WO 9006371, U.S. Pat. No. 4,935,343, EP 364778, EP 267611 and EP 220063;
- 20 • IL-1 receptor accessory proteins and antibodies thereto (e.g., WO 96/23067);
- Inhibitors of interleukin-1 β converting enzyme (ICE) or caspase I, which can be used to inhibit IL-1 beta production and secretion;
- 25 • Interleukin-1 β protease inhibitors;
- Other compounds and proteins which block *in vivo* synthesis or extracellular release of IL-1.

Exemplary IL-1 inhibitors are disclosed in the following references:

30 US Pat. Nos. 5747444; 5359032; 5608035; 5843905; 5359032; 5866576; 5869660; 5869315; 5872095; 5955480;

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International (WO) patent applications 98/21957,
96/09323, 91/17184, 96/40907, 98/32733, 98/42325, 98/44940,
98/47892, 98/56377, 99/03837, 99/06426, 99/06042, 91/17249,
98/32733, 98/17661, 97/08174, 95/34326, 99/36426, and
5 99/36415;

European (EP) patent applications 534978 and 894795; and
French patent application FR 2762514;

Interleukin-1 receptor antagonist (IL-1ra) is a human protein that acts as a natural inhibitor of interleukin-1.
10 Preferred receptor antagonists (including IL-1ra and variants and derivatives thereof), as well as methods of making and using thereof, are described in U.S. Patent No. 5,075,222; WO 91/08285; WO 91/17184; AU 9173636; WO 92/16221; WO93/21946; WO 94/06457; WO 94/21275; FR 2706772; WO 94/21235; DE 4219626, WO 15 94/20517; WO 96/22793; WO 97/28828; and WO 99/36541. The proteins include glycosylated as well as non-glycosylated IL-1 receptor antagonists.

Specifically, three exemplary forms of IL-1ra and variants thereof are disclosed and described in the 5,075,222 patent. The first of these, called "IL-1 α " in the 5,075,222 patent, is characterized as a 22-23 kD molecule on SDS-PAGE with an approximate isoelectric point of 4.8, eluting from a MonoQ FPLC column at around 52 mM NaCl in Tris buffer, pH 7.6. The second, IL-1ra β , is characterized as a 22-23 kD protein, 25 eluting from a MonoQ column at 48 mM NaCl. Both IL-1ra α and IL-1ra β are glycosylated. The third, IL-1rax, is characterized as a 20 kD protein, eluting from a MonoQ column at 48 mM NaCl, and is non-glycosylated. U.S. patent No. 5,075,222 also discloses methods for isolating the genes 30 responsible for coding the inhibitors, cloning the gene in suitable vectors and cell types, and expressing the gene to produce the inhibitors.

Those skilled in the art will recognize that many combinations of deletions, insertions, and substitutions

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(individually or collectively "variant(s)" herein) can be made within the amino acid sequences of IL-1ra, provided that the resulting molecule is biologically active (e.g., possesses the ability to affect one or more of the diseases and disorders 5 such as those recited herein.)

As contemplated by the present invention, an agonist or antagonist of the IL-17 receptor like polypeptide (including, but not limited to, anti-IL-17 receptor like selective binding agents [such as antibodies], ligands to the IL-17 RECEPTOR 10 like receptor, soluble IL-17 receptor like polypeptides, small molecules, and antisense oligonucleotides, or an IL-17 receptor like polypeptide itself) may be administered as an adjunct to other therapy and also with other pharmaceutical compositions suitable for the indication being treated. An 15 agonist or antagonist of the IL-17 receptor like polypeptide, and/or an IL-17 receptor like polypeptide itself and any of one or more additional therapies or pharmaceutical formulations may be administered separately, sequentially, or simultaneously.

20

In a specific embodiment, the present invention is directed to the use of an agonist or antagonist of the IL-17 receptor like polypeptide, and/or an IL-17 receptor like polypeptide itself in combination (pre-treatment, post- 25 treatment, or concurrent treatment) with any of one or more TNF inhibitors for the treatment or prevention of the diseases and disorders recited herein.

Such TNF inhibitors include compounds and proteins which block *in vivo* synthesis or extracellular release of TNF. In a 30 specific embodiment, the present invention is directed to the use of an agonist or antagonist of the IL-17 receptor like polypeptide, and/or an IL-17 receptor like polypeptide itself in combination (pre-treatment, post-treatment, or concurrent treatment) with any of one or more of the following TNF

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inhibitors: TNF binding proteins (soluble TNF receptor type-I and soluble TNF receptor type-II ("sTNFRs"), as defined herein), anti-TNF antibodies, granulocyte colony stimulating factor; thalidomide; BN 50730; tenidap; E 5531; tiapafant PCA
5 4248; nimesulide; panavir; rolipram; RP 73401; peptide T; MDL 201,449A; (1R,3S)-Cis-1-[9-(2,6-diaminopurinyl)]-3-hydroxy-4-cyclopentene hydrochloride; (1R,3R)-trans-1-(9-(2,6-diamino)purine]-3-acetoxycyclopentane; (1R,3R)-trans-1-[9-adenyl]-3-azidocyclopentane hydrochloride and (1R,3R)-trans-1-
10 (6-hydroxy-purin-9-yl)-3-azidocyclo-pentane. TNF binding proteins are disclosed in the art (EP 308 378, EP 422 339, GB 2 218 101, EP 393 438, WO 90/13575, EP 398 327, EP 412 486, WO 91/03553, EP 418 014, JP 127,800/1991, EP 433 900, U.S. Patent No. 5,136,021, GB 2 246 569, EP 464 533, WO 92/01002,
15 WO 92/13095, WO 92/16221, EP 512 528, EP 526 905, WO 93/07863, EP 568 928, WO 93/21946, WO 93/19777, EP 417 563, WO 94/06476, and PCT International Application No. PCT/US97/12244).

For example, EP 393 438 and EP 422 339 teach the amino acid and nucleic acid sequences of a soluble TNF receptor type
20 I (also known as "sTNFR-I" or "30kDa TNF inhibitor") and a soluble TNF receptor type II (also known as "sTNFR-II" or "40kDa TNF inhibitor"), collectively termed "sTNFRs", as well as modified forms thereof (e.g., fragments, functional derivatives and variants). EP 393 438 and EP 422 339 also disclose methods for isolating the genes responsible for coding the inhibitors, cloning the gene in suitable vectors and cell types and expressing the gene to produce the inhibitors. Additionally, polyvalent forms (i.e., molecules comprising more than one active moiety) of sTNFR-I and sTNFR-II have also been disclosed. In one embodiment, the polyvalent form may be constructed by chemically coupling at least one TNF inhibitor and another moiety with any clinically acceptable linker, for example polyethylene glycol (WO 92/16221 and WO 95/34326), by a peptide linker (Neve et
30 al. (1996), *Cytokine*, 8(5):365-370, by chemically coupling to
35

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biotin and then binding to avidin (WO 91/03553) and, finally, by combining chimeric antibody molecules (U.S. Patent 5,116,964, WO 89/09622, WO 91/16437 and EP 315062.

Anti-TNF antibodies include MAK 195F Fab antibody (Holler et al. (1993), 1st International Symposium on Cytokines in Bone Marrow Transplantation, 147); CDP 571 anti-TNF monoclonal antibody (Rankin et al. (1995), *British Journal of Rheumatology*, 34:334-342); BAY X 1351 murine anti-tumor necrosis factor monoclonal antibody (Kieft et al. (1995), 10 7th European Congress of Clinical Microbiology and Infectious Diseases, page 9); CenTNF cA2 anti-TNF monoclonal antibody (Elliott et al. (1994), *Lancet*, 344:1125-1127 and Elliott et al. (1994), *Lancet*, 344:1105-1110).

In a specific embodiment, the present invention is directed to the use of agonist or antagonist of the IL-17 receptor like polypeptide, and/or an IL-17 receptor like polypeptide itself in combination (pretreatment, post-treatment, or concurrent treatment) with secreted or soluble human fas antigen or recombinant versions thereof (WO 96/20206 and Mountz et al., *J. Immunology*, 155:4829-4837; and EP 510 691. WO 96/20206 discloses secreted human fas antigen (native and recombinant, including an Ig fusion protein), methods for isolating the genes responsible for coding the soluble recombinant human fas antigen, methods for cloning the gene in suitable vectors and cell types, and methods for expressing the gene to produce the inhibitors. EP 510 691 describes DNAs coding for human fas antigen, including soluble fas antigen, vectors expressing for said DNAs and transformants transfected with the vector. When administered parenterally, doses of a 20 secreted or soluble fas antigen fusion protein each are generally from about 1 micrograms/kg to about 100 micrograms/kg.

Treatment of the diseases and disorders recited herein, including acute and chronic inflammation such as rheumatic

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diseases, commonly includes the use of first line drugs for control of pain and inflammation; these drugs are classified as non-steroidal, anti-inflammatory drugs (NSAIDs). Secondary treatments include corticosteroids, slow acting antirheumatic 5 drugs (SAARDs), or disease modifying (DM) drugs. Information regarding the following compounds can be found in The Merck Manual of Diagnosis and Therapy, Sixteenth Edition, Merck, Sharp & Dohme Research Laboratories, Merck & Co., Rahway, NJ (1992) and in Pharmaprojects, PJB Publications Ltd.

10 In a specific embodiment, the present invention is directed to the use of an agonist or antagonist of the IL-17 receptor like polypeptide, and/or an IL-17 receptor like polypeptide itself and any of one or more NSAIDs for the treatment of the diseases and disorders recited herein, 15 including acute and chronic inflammation such as rheumatic diseases; and graft versus host disease. NSAIDs owe their anti-inflammatory action, at least in part, to the inhibition of prostaglandin synthesis (Goodman and Gilman in "The Pharmacological Basis of Therapeutics," MacMillan 7th Edition 20 (1985)). NSAIDs can be characterized into at least nine groups: (1) salicylic acid derivatives; (2) propionic acid derivatives; (3) acetic acid derivatives; (4) fenamic acid derivatives; (5) carboxylic acid derivatives; (6) butyric acid derivatives; (7) oxicams; (8) pyrazoles and (9) pyrazolones.

25 In another specific embodiment, the present invention is directed to the use of an agonist or antagonist of the IL-17 receptor like polypeptide, and/or an IL-17 receptor like polypeptide itself in combination (pretreatment, post-treatment, or concurrent treatment) with any of one or more 30 salicylic acid derivatives, prodrug esters or pharmaceutically acceptable salts thereof. Such salicylic acid derivatives, prodrug esters and pharmaceutically acceptable salts thereof comprise: acetaminosalol, aloxiprin, aspirin, benorylate, bromosaligenin, calcium acetylsalicylate, choline magnesium 35 trisalicylate, magnesium salicylate, choline salicylate,

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diflusinal, etersalate, fendosal, gentisic acid, glycol salicylate, imidazole salicylate, lysine acetylsalicylate, mesalamine, morpholine salicylate, 1-naphthyl salicylate, olsalazine, parsalmide, phenyl acetylsalicylate, phenyl 5 salicylate, salacetamide, salicylamide O-acetic acid, salsalate, sodium salicylate and sulfasalazine. Structurally related salicylic acid derivatives having similar analgesic and anti-inflammatory properties are also intended to be encompassed by this group.

10 In an additional specific embodiment, the present invention is directed to the use of an agonist or antagonist of the IL-17 receptor like polypeptide, and/or an IL-17 receptor like polypeptide itself in combination (pretreatment, post-treatment, or concurrent treatment) with any of one or 15 more propionic acid derivatives, prodrug esters or pharmaceutically acceptable salts thereof. The propionic acid derivatives, prodrug esters, and pharmaceutically acceptable salts thereof comprise: alminoprofen, benoxaprofen, bucloxic acid, carprofen, dexindoprofen, fenoprofen, flunoxaprofen, 20 fluprofen, flurbiprofen, furclopofen, ibuprofen, ibuprofen aluminum, ibuproxam, indoprofen, isoprofen, ketoprofen, loxoprofen, mioprofen, naproxen, naproxen sodium, oxaprozin, piketoprofen, pimeprofen, pirprofen, pranoprofen, protizinic acid, pyridoxiprofen, suprofen, tiaprofenic acid and 25 tioxaprofen. Structurally related propionic acid derivatives having similar analgesic and anti-inflammatory properties are also intended to be encompassed by this group.

In yet another specific embodiment, the present invention is directed to the use of an agonist or antagonist of the IL- 30 17 receptor like polypeptide, and/or an IL-17 receptor like polypeptide itself in combination (pretreatment, post-treatment, or concurrent treatment) with any of one or more acetic acid derivatives, prodrug esters or pharmaceutically acceptable salts thereof. The acetic acid derivatives, 35 prodrug esters, and pharmaceutically acceptable salts thereof

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comprise: acemetacin, alclofenac, amfenac, bufexamac,
cinmetacin, clopirac, delmetacin, diclofenac potassium,
diclofenac sodium, etodolac, felbinac, fenclofenac, fenclorac,
fenclozic acid; fentiazac, furofenac, glucametacin, ibufenac,
5 indomethacin, isofezolac, isoxepac, lonazolac, metiazinic
acid, oxametacin, oxpipac, pimetacin, proglumetacin, sulindac,
talmetacin, tiaramide, tiopinac, tolmetin, tolmetin sodium,
zidometacin and zomepirac. Structurally related acetic acid
derivatives having similar analgesic and anti-inflammatory
10 properties are also intended to be encompassed by this group.

In another specific embodiment, the present invention is directed to the use of an agonist or antagonist of the IL-17 receptor like polypeptide, and/or an IL-17 receptor like polypeptide itself in combination (pretreatment, post-treatment, or concurrent treatment) with any of one or more fenamic acid derivatives, prodrug esters or pharmaceutically acceptable salts thereof. The fenamic acid derivatives, prodrug esters and pharmaceutically acceptable salts thereof comprise: enfenamic acid, etofenamate, flufenamic acid,
20 isonixin, meclofenamic acid, meclofenamate sodium, medofenamic acid, mefenamic acid, niflumic acid, talniflumate, terofenamate, tolfenamic acid and ufenamate. Structurally related fenamic acid derivatives having similar analgesic and anti-inflammatory properties are also intended to be
25 encompassed by this group.

In an additional specific embodiment, the present invention is directed to the use of an agonist or antagonist of the IL-17 receptor like polypeptide, and/or an IL-17 receptor like polypeptide itself in combination (pretreatment, post-treatment, or concurrent treatment) with any of one or more carboxylic acid derivatives, prodrug esters or pharmaceutically acceptable salts thereof. The carboxylic acid derivatives, prodrug esters, and pharmaceutically acceptable salts thereof which can be used comprise: clidanac,
30 diflunisal, flufenisal, inoridine, ketorolac and tinoridine.

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Structurally related carboxylic acid derivatives having similar analgesic and anti-inflammatory properties are also intended to be encompassed by this group.

In yet another specific embodiment, the present invention 5 is directed to the use of an agonist or antagonist of the IL-17 receptor like polypeptide, and/or an IL-17 receptor like polypeptide itself in combination (pretreatment, post-treatment, or concurrent treatment) with any of one or more butyric acid derivatives, prodrug esters or pharmaceutically 10 acceptable salts thereof. The butyric acid derivatives, prodrug esters, and pharmaceutically acceptable salts thereof comprise: bumadizon, butibufen, fenbufen and xenbucin. Structurally related butyric acid derivatives having similar analgesic and anti-inflammatory properties are also intended 15 to be encompassed by this group.

In another specific embodiment, the present invention is directed to the use of an agonist or antagonist of the IL-17 receptor like polypeptide, and/or an IL-17 receptor like polypeptide itself in combination (pretreatment, post-treatment, or concurrent treatment) with any of one or more 20 oxicams, prodrug esters, or pharmaceutically acceptable salts thereof. The oxicams, prodrug esters, and pharmaceutically acceptable salts thereof comprise: droxicam, enolicam, isoxicam, piroxicam, sudoxicam, tenoxicam and 4-hydroxyl-1,2-benzothiazine-1,1-dioxide-4-(N-phenyl)-carboxamide. Structurally 25 related oxicams having similar analgesic and anti-inflammatory properties are also intended to be encompassed by this group.

In still another specific embodiment, the present 30 invention is directed to the use of an agonist or antagonist of the IL-17 receptor like polypeptide, and/or an IL-17 receptor like polypeptide itself in combination (pretreatment, post-treatment, or concurrent treatment) with any of one or more pyrazoles, prodrug esters, or pharmaceutically acceptable

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salts thereof. The pyrazoles, prodrug esters, and pharmaceutically acceptable salts thereof which may be used comprise: difenamizole and spirizole. Structurally related pyrazoles having similar analgesic and anti-inflammatory properties are also intended to be encompassed by this group.

In an additional specific embodiment, the present invention is directed to the use of an agonist or antagonist of the IL-17 receptor like polypeptide, and/or an IL-17 receptor like polypeptide itself in combination (pretreatment, post-treatment or, concurrent treatment) with any of one or more pyrazolones, prodrug esters, or pharmaceutically acceptable salts thereof. The pyrazolones, prodrug esters and pharmaceutically acceptable salts thereof which may be used comprise: apazone, azapropazone, benzpiperylon, feprazone, mofebutazone, morazone, oxyphenbutazone, phenylbutazone, pipebuzone, propylphenazone, ramifenazone, suxibuzone and thiazolinobutazone. Structurally related pyrazolones having similar analgesic and anti-inflammatory properties are also intended to be encompassed by this group.

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In another specific embodiment, the present invention is directed to the use of an agonist or antagonist of the IL-17 receptor like polypeptide, and/or an IL-17 receptor like polypeptide itself in combination (pretreatment, post-treatment, or concurrent treatment) with any of one or more of the following NSAIDs: ϵ -acetamidocaproic acid, S-adenosylmethionine, 3-amino-4-hydroxybutyric acid, amixetrine, anitrazafen, antrafenine, bendazac, bendazac lysinate, benzydamine, beprozin, broperamide, bucolome, bufezolac, ciproquazone, cloximate, dazidamine, deboxamet, detomidine, difenpiramide, difenpyramide, difisalamine, ditazol, emorfazole mesylate, fenflumizole, floctafenine, flumizole, flunixin, fluproquazone, fopirtoline, fosfosal, guaimesal, guaiazolene, isonixirn, lefetamine HCl,

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leflunomide, lofemizole, lotifazole, lysin clonixinate,
meseclazone, nabumetone, nictindole, nimesulide, orgotein,
orpanoxin, oxaceprol, oxapadol, paranyline, perisoxal,
perisoxal citrate, pifoxime, piproxen, pirazolac, pirfenidone,
5 proquazone, proxazole, thielavin B, tiflamizole, timegadine,
tolectin, tolpadol, tryptamid and those designated by company
code number such as 480156S, AA861, AD1590, AFP802, AFP860,
AI77B, AP504, AU8001, BPPC, BW540C, CHINOIN 127, CN100, EB382,
EL508, F1044, FK-506, GV3658, ITF182, KCNTEI6090, KME4,
10 LA2851, MR714, MR897, MY309, ONO3144, PR823, PV102, PV108,
R830, RS2131, SCR152, SH440, SIR133, SPAS510, SQ27239, ST281,
SY6001, TA60, TAI-901 (4-benzoyl-1-indancarboxylic acid),
TVX2706, U60257, UR2301 and WY41770. Structurally related
15 NSAIDs having similar analgesic and anti-inflammatory
properties to the NSAIDs are also intended to be encompassed
by this group.

In still another specific embodiment, the present invention is directed to the use of an agonist or antagonist of the IL-17 receptor like polypeptide, and/or an IL-17 receptor like polypeptide itself in combination (pretreatment, post-treatment or concurrent treatment) with any of one or more corticosteroids, prodrug esters or pharmaceutically acceptable salts thereof for the treatment of the diseases and disorders recited herein, including acute and chronic inflammation such as rheumatic diseases, graft versus host disease and multiple sclerosis. Corticosteroids, prodrug esters and pharmaceutically acceptable salts thereof include hydrocortisone and compounds which are derived from hydrocortisone, such as 21-acetoxypregnенolone, alclomerasone, 20 algestone, amcinonide, beclomethasone, betamethasone, betamethasone valerate, budesonide, chloroprednisone, clobetasol, clobetasol propionate, clobetasone, clobetasone butyrate, clocortolone, cloprednol, corticosterone, cortisone, cortivazol, deflazacon, desonide, desoximerasone, 30 dexamethasone, diflorasone, diflucortolone, difluprednate,

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enoxolone, fluazacort, flucloronide, flumethasone,
flumethasone pivalate, flucinolone acetonide, flunisolide,
fluocinonide, fluorocinolone acetonide, fluocortin butyl,
fluocortolone, fluocortolone hexanoate, diflucortolone
5 valerate, fluorometholone, fluperolone acetate, fluprednidene
acetate, fluprednisolone, flurandenolide, formocortal,
halcinonide, halometasone, halopredone acetate,
hydrocortamate, hydrocortisone, hydrocortisone acetate,
hydrocortisone butyrate, hydrocortisone phosphate,
10 hydrocortisone 21-sodium succinate, hydrocortisone tebutate,
mazipredone, medrysone, meprednisone, methylprednisolone,
mometasone furoate, paramethasone, prednicarbate,
prednisolone, prednisolone 21-diedryaminoacetate, prednisolone
sodium phosphate, prednisolone sodium succinate, prednisolone
15 sodium 21-m-sulfobenzoate, prednisolone sodium 21-
stearoglycolate, prednisolone tebutate, prednisolone 21-
trimethylacetate, prednisone, prednival, prednylidene,
prednylidene 21-diethylaminoacetate, tixocortol,
triamcinolone, triamcinolone acetonide, triamcinolone
20 benetonide and triamcinolone hexacetonide. Structurally
related corticosteroids having similar analgesic and anti-
inflammatory properties are also intended to be encompassed by
this group.

In another specific embodiment, the present invention is
25 directed to the use of an agonist or antagonist of the IL-17
receptor like polypeptide, and/or an IL-17 receptor like
polypeptide itself in combination (pretreatment, post-
treatment, or concurrent treatment) with any of one or more
slow-acting antirheumatic drugs (SAARDs) or disease modifying
30 antirheumatic drugs (DMARDs), prodrug esters, or
pharmaceutically acceptable salts thereof for the treatment of
the diseases and disorders recited herein, including acute and
chronic inflammation such as rheumatic diseases, graft versus
host disease and multiple sclerosis. SAARDs or DMARDs,
35 prodrug esters and pharmaceutically acceptable salts thereof

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comprise: allocupreide sodium, auranofin, aurothioglucose,
aurothioglycanide, azathioprine, brequinar sodium,
bucillamine, calcium 3-aurothio-2-propanol-1-sulfonate,
chlorambucil, chloroquine, clobuzarit, cuproxoline,
5 cyclophosphamide, cyclosporin, dapsone, 15-deoxyspergualin,
diacerein, glucosamine, gold salts (e.g., cycloquine gold
salt, gold sodium thiomalate, gold sodium thiosulfate),
hydroxychloroquine, hydroxychloroquine sulfate, hydroxyurea,
kebuzone, levamisole, lobenzarit, melittin, 6-mercaptopurine,
10 methotrexate, mizoribine, mycophenolate mofetil, myoral,
nitrogen mustard, D-penicillamine, pyridinol imidazoles such
as SKNF86002 and SB203580, rapamycin, thiols, thymopoietin and
vincristine. Structurally related SAARDs or DMARDs having
similar analgesic and anti-inflammatory properties are also
15 intended to be encompassed by this group.

In another specific embodiment, the present invention is directed to the use of an agonist or antagonist of the IL-17 receptor like polypeptide, and/or an IL-17 receptor like polypeptide itself in combination (pretreatment, post-treatment, or concurrent treatment) with any of one or more COX2 inhibitors, prodrug esters or pharmaceutically acceptable salts thereof for the treatment of the diseases and disorders recited herein; including acute and chronic inflammation. Examples of COX2 inhibitors, prodrug esters or 25 pharmaceutically acceptable salts thereof include, for example, celecoxib. Structurally related COX2 inhibitors having similar analgesic and anti-inflammatory properties are also intended to be encompassed by this group.

In still another specific embodiment, the present invention is directed to the use of an agonist or antagonist of the IL-17 receptor like polypeptide, and/or an IL-17 receptor like polypeptide itself in combination (pretreatment, post-treatment, or concurrent treatment) with any of one or more antimicrobials, prodrug esters or pharmaceutically acceptable salts thereof for the treatment of the diseases and

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disorders recited herein, including acute and chronic inflammation. Antimicrobials include, for example, the broad classes of penicillins, cephalosporins and other beta-lactams, aminoglycosides, azoles, quinolones, macrolides, rifamycins, 5 tetracyclines, sulfonamides, lincosamides and polymyxins. The penicillins include, but are not limited to penicillin G, penicillin V, methicillin, nafcillin, oxacillin, cloxacillin, dicloxacillin, floxacillin, ampicillin, ampicillin/sulbactam, amoxicillin, amoxicillin/clavulanate, hetacillin, cyclacillin, 10 bacampicillin, carbenicillin, carbenicillin indanyl, ticarcillin, ticarcillin/clavulanate, azlocillin, mezlocillin, peperacillin, and mecillinam. The cephalosporins and other beta-lactams include, but are not limited to cephalothin, cephapirin, cephalexin, cephradine, cefazolin, cefadroxil, 15 cefaclor, cefamandole, cefotetan, cefoxitin, ceruroxime, cefonicid, ceforadine, cefixime, cefotaxime, moxalactam, ceftizoxime, ceftriaxone, cephoperazone, ceftazidime, imipenem and aztreonam. The aminoglycosides include, but are not limited to streptomycin, gentamicin, tobramycin, amikacin, 20 netilmicin, kanamycin and neomycin. The azoles include, but are not limited to fluconazole. The quinolones include, but are not limited to nalidixic acid, norfloxacin, enoxacin, ciprofloxacin, ofloxacin, sparfloxacin and temafloxacin. The macrolides include, but are not limited to erythromycin, 25 spiramycin and azithromycin. The rifamycins include, but are not limited to rifampin. The tetracyclines include, but are not limited to spirocycline, chlortetracycline, clomocycline, demeclocycline, doxycycline, guamecycline, lymecycline, meclocycline, methacycline, minocycline, oxytetracycline, 30 penimepicycline, pipacycline, rolitetracycline, sancycline, senociclin and tetracycline. The sulfonamides include, but are not limited to sulfanilamide, sulfamethoxazole, sulfacetamide, sulfadiazine, sulfisoxazole and co-trimoxazole (trimethoprim/sulfamethoxazole). The lincosamides include, 35 but are not limited to clindamycin and lincomycin. The

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polymyxins (polypeptides) include, but are not limited to polymyxin B and colistin.

IL-17 Receptor Like Compositions and Administration

Therapeutic compositions are within the scope of the present invention. Such IL-17 receptor like pharmaceutical compositions may comprise a therapeutically effective amount of an IL-17 receptor like polypeptide or an IL-17 receptor like nucleic acid molecule in admixture with a pharmaceutically or physiologically acceptable formulation agent selected for suitability with the mode of administration. Pharmaceutical compositions may comprise a therapeutically effective amount of one or more IL-17 receptor like selective binding agents in admixture with a pharmaceutically or physiologically acceptable formulation agent selected for suitability with the mode of administration.

Acceptable formulation materials preferably are nontoxic to recipients at the dosages and concentrations employed.

The pharmaceutical composition may contain formulation materials for modifying, maintaining or preserving, for example, the pH, osmolarity, viscosity, clarity, color, isotonicity, odor, sterility, stability, rate of dissolution or release, adsorption or penetration of the composition. Suitable formulation materials include, but are not limited to, amino acids (such as glycine, glutamine, asparagine, arginine or lysine), antimicrobials, antioxidants (such as ascorbic acid, sodium sulfite or sodium hydrogen-sulfite), buffers (such as borate, bicarbonate, Tris-HCl, citrates, phosphates, other organic acids), bulking agents (such as mannitol or glycine), chelating agents (such as ethylenediamine tetraacetic acid (EDTA)), complexing agents

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(such as caffeine, polyvinylpyrrolidone, beta-cyclodextrin or hydroxypropyl-beta-cyclodextrin), fillers, monosaccharides, disaccharides, and other carbohydrates (such as glucose, mannose, or dextrins), proteins (such as serum albumin, 5 gelatin or immunoglobulins), coloring, flavoring and diluting agents, emulsifying agents, hydrophilic polymers (such as polyvinylpyrrolidone), low molecular weight polypeptides, salt-forming counterions (such as sodium), preservatives (such as benzalkonium chloride, benzoic acid, salicylic acid, 10 thimerosal, phenethyl alcohol, methylparaben, propylparaben, chlorhexidine, sorbic acid or hydrogen peroxide), solvents (such as glycerin, propylene glycol or polyethylene glycol), sugar alcohols (such as mannitol or sorbitol), suspending agents, surfactants or wetting agents (such as pluronic, PEG, 15 sorbitan esters, polysorbates such as polysorbate 20, polysorbate 80, triton, tromethamine, lecithin, cholesterol, tyloxapal), stability enhancing agents (sucrose or sorbitol), tonicity enhancing agents (such as alkali metal halides (preferably sodium or potassium chloride), mannitol sorbitol), 20 delivery vehicles, diluents, excipients and/or pharmaceutical adjuvants. See *Remington's Pharmaceutical Sciences*, 18th Ed., A.R. Gennaro, ed., Mack Publishing Company (1990).

The optimal pharmaceutical composition will be determined by one skilled in the art depending upon, for example, the 25 intended route of administration, delivery format, and desired dosage. See for example, *Remington's Pharmaceutical Sciences*, supra. Such compositions may influence the physical state, stability, rate of *in vivo* release, and rate of *in vivo* clearance of the IL-17 receptor like molecule.

30 The primary vehicle or carrier in a pharmaceutical composition may be either aqueous or non-aqueous in nature. For example, a suitable vehicle or carrier may be water for injection, physiological saline solution, or artificial cerebrospinal fluid, possibly supplemented with other 35 materials common in compositions for parenteral

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administration. Neutral buffered saline or saline mixed with serum albumin are further exemplary vehicles. Other exemplary pharmaceutical compositions comprise Tris buffer of about pH 7.0-8.5, or acetate buffer of about pH 4.0-5.5, which may 5 further include sorbitol or a suitable substitute therefor. In one embodiment of the present invention, IL-17 receptor like polypeptide compositions may be prepared for storage by mixing the selected composition having the desired degree of purity with optional formulation agents (*Remington's* 10 *Pharmaceutical Sciences, supra*) in the form of a lyophilized cake or an aqueous solution. Further, the IL-17 receptor like polypeptide product may be formulated as a lyophilizate using appropriate excipients such as sucrose.

The IL-17 receptor like pharmaceutical compositions can 15 be selected for parenteral delivery. Alternatively, the compositions may be selected for inhalation or for delivery through the digestive tract, such as orally. The preparation of such pharmaceutically acceptable compositions is within the skill of the art.

20 The formulation components are present in concentrations that are acceptable to the site of administration. For example, buffers are used to maintain the composition at physiological pH or at slightly lower pH, typically within a pH range of from about 5 to about 8.

25 When parenteral administration is contemplated, the therapeutic compositions for use in this invention may be in the form of a pyrogen-free, parenterally acceptable aqueous solution comprising the desired IL-17 receptor like molecule in a pharmaceutically acceptable vehicle. A particularly 30 suitable vehicle for parenteral injection is sterile distilled water in which a IL-17 receptor like molecule is formulated as a sterile, isotonic solution, properly preserved. Yet another preparation can involve the formulation of the desired molecule with an agent, such as injectable microspheres, bio- 35 erodible particles, polymeric compounds (polylactic acid,

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polyglycolic acid), or beads, or liposomes, that provides for the controlled or sustained release of the product which may then be delivered as a depot injection. Hyaluronic acid may also be used, and this may have the effect of promoting 5 sustained duration in the circulation. Other suitable means for the introduction of the desired molecule include implantable drug delivery devices.

Pharmaceutical compositions such as (1) slow-release formulations, (2) inhalant mists, or (3) orally active 10 formulations are also envisioned. The IL-17 receptor like molecule pharmaceutical composition generally is formulated for parenteral administration. Such parenterally administered therapeutic compositions are typically in the form of a pyrogen-free, parenterally acceptable aqueous solution 15 comprising the desired IL-17 receptor like molecule in a pharmaceutically acceptable vehicle. The IL-17 receptor like molecule pharmaceutical compositions also may include particulate preparations of polymeric compounds such as polylactic acid, polyglycolic acid, etc. or the introduction 20 of the molecule into liposomes. Hyaluronic acid may also be used, and this may have the effect of promoting sustained duration in the circulation

In one embodiment, a pharmaceutical composition may be formulated for inhalation. For example, an IL-17 receptor 25 like molecule may be formulated as a dry powder for inhalation. IL-17 receptor like polypeptide or IL-17 receptor like nucleic acid molecule inhalation solutions may also be formulated with a propellant for aerosol delivery. In yet another embodiment, solutions may be nebulized. Pulmonary 30 administration is further described in PCT application no. PCT/US94/001875, which describes pulmonary delivery of chemically modified proteins.

It is also contemplated that certain formulations may be administered orally. In one embodiment of the present 35 invention, IL-17 receptor like molecules which are

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administered in this fashion can be formulated with or without those carriers customarily used in the compounding of solid dosage forms such as tablets and capsules. For example, a capsule may be designed to release the active portion of the
5 formulation at the point in the gastrointestinal tract when bioavailability is maximized and pre-systemic degradation is minimized. Additional agents can be included to facilitate absorption of the IL-17 receptor like molecule. Diluents, flavorings, low melting point waxes, vegetable oils,
10 lubricants, suspending agents, tablet disintegrating agents, and binders may also be employed.

Another pharmaceutical composition may involve an effective quantity of IL-17 receptor like molecules in a mixture with non-toxic excipients which are suitable for the
15 manufacture of tablets. By dissolving the tablets in sterile water, or other appropriate vehicle, solutions can be prepared in unit dose form. Suitable excipients include, but are not limited to, inert diluents, such as calcium carbonate, sodium carbonate or bicarbonate, lactose, or calcium phosphate; or
20 binding agents, such as starch, gelatin, or acacia; or lubricating agents such as magnesium stearate, stearic acid, or talc.

Additional IL-17 receptor like pharmaceutical compositions will be evident to those skilled in the art,
25 including formulations involving IL-17 receptor like polypeptides in sustained- or controlled-delivery formulations. Techniques for formulating a variety of other sustained- or controlled-delivery means, such as liposome carriers, bio-erodible microparticles or porous beads and
30 depot injections, are also known to those skilled in the art. See for example, PCT/US93/00829 which describes controlled release of porous polymeric microparticles for the delivery of pharmaceutical compositions. Additional examples of sustained-release preparations include semipermeable polymer
35 matrices in the form of shaped articles, e.g. films, or

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microcapsules. Sustained release matrices may include polyesters, hydrogels, polylactides (U.S. 3,773,919, EP 58,481), copolymers of L-glutamic acid and gamma ethyl-L-glutamate (Sidman et al., *Biopolymers*, 22:547-556 (1983)),
5 poly (2-hydroxyethyl-methacrylate) (Langer et al., *J. Biomed. Mater. Res.*, 15:167-277 (1981) and Langer, *Chem. Tech.*, 12:98-105 (1982)), ethylene vinyl acetate (Langer et al., *supra*) or poly-D(-)-3-hydroxybutyric acid (EP 133,988). Sustained-release compositions also may include liposomes, which can be
10 prepared by any of several methods known in the art. See e.g., Eppstein et al., *Proc. Natl. Acad. Sci. USA*, 82:3688-3692 (1985); EP 36,676; EP 88,046; EP 143,949.

The IL-17 receptor like pharmaceutical composition to be used for *in vivo* administration typically must be sterile.
15 This may be accomplished by filtration through sterile filtration membranes. Where the composition is lyophilized, sterilization using these methods may be conducted either prior to, or following, lyophilization and reconstitution. The composition for parenteral administration may be stored in
20 lyophilized form or in solution. In addition, parenteral compositions generally are placed into a container having a sterile access port, for example, an intravenous solution bag or vial having a stopper pierceable by a hypodermic injection needle.

Once the pharmaceutical composition has been formulated, it may be stored in sterile vials as a solution, suspension, gel, emulsion, solid, or a dehydrated or lyophilized powder. Such formulations may be stored either in a ready-to-use form or in a form (e.g., lyophilized) requiring reconstitution
30 prior to administration.

In a specific embodiment, the present invention is directed to kits for producing a single-dose administration unit. The kits may each contain both a first container having a dried protein and a second container having an aqueous

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formulation. Also included within the scope of this invention are kits containing single and multi-chambered pre-filled syringes (e.g., liquid syringes and lyosyringes).

An effective amount of an IL-17 receptor like pharmaceutical composition to be employed therapeutically will depend, for example, upon the therapeutic context and objectives. One skilled in the art will appreciate that the appropriate dosage levels for treatment will thus vary depending, in part, upon the molecule delivered, the indication for which the IL-17 receptor like molecule is being used, the route of administration, and the size (body weight, body surface or organ size) and condition (the age and general health) of the patient. Accordingly, the clinician may titer the dosage and modify the route of administration to obtain the optimal therapeutic effect. A typical dosage may range from about 0.1 $\mu\text{g}/\text{kg}$ to up to about 100 mg/kg or more, depending on the factors mentioned above. In other embodiments, the dosage may range from 0.1 $\mu\text{g}/\text{kg}$ up to about 100 mg/kg ; or 1 $\mu\text{g}/\text{kg}$ up to about 100 mg/kg ; or 5 $\mu\text{g}/\text{kg}$ up to about 100 mg/kg .

The frequency of dosing will depend upon the pharmacokinetic parameters of the IL-17 receptor like molecule in the formulation used. Typically, a clinician will administer the composition until a dosage is reached that achieves the desired effect. The composition may therefore be administered as a single dose, or as two or more doses (which may or may not contain the same amount of the desired molecule) over time, or as a continuous infusion via implantation device or catheter. Further refinement of the appropriate dosage is routinely made by those of ordinary skill in the art and is within the ambit of tasks routinely performed by them. Appropriate dosages may be ascertained through use of appropriate dose-response data.

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The route of administration of the pharmaceutical composition is in accord with known methods, e.g. oral, inhalation, injection or infusion by intravenous, intraperitoneal, intracerebral (intra-parenchymal),
5 intracerebroventricular, intramuscular, intra-ocular, intraarterial, intraportal, or intralesional routes, or by sustained release systems or implantation device. Where desired, the compositions may be administered continuously by infusion, by bolus injection or continuously by infusion, or
10 by implantation device.

Alternatively or additionally, the composition may be administered locally via implantation into the affected area of a membrane, sponge, or other appropriate material on to which the desired molecule has been absorbed or encapsulated.
15 Where an implantation device is used, the device may be implanted into any suitable tissue or organ, and delivery of the desired molecule may be via diffusion, timed release bolus, or continuous administration or via catheter using continuous infusion.

20 It will further be appreciated that the IL-17 receptor like polypeptides, including fragments, variants, and derivatives, may be employed alone, together, or in combination with other polypeptides and pharmaceutical compositions. For example, the IL-17 receptor like
25 polypeptides may be used in combination with cytokines, growth factors, antibiotics, anti-inflammatories, and/or chemotherapeutic agents as is appropriate for the indication being treated.

In some cases, it may be desirable to use IL-17 receptor
30 like pharmaceutical compositions in an ex vivo manner. In such instances, cells, tissues, or organs that have been removed from the patient are exposed to IL-17 receptor like pharmaceutical compositions after which the cells, tissues

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and/or organs are subsequently implanted back into the patient.

In other cases, an IL-17 receptor like polypeptide can be delivered by implanting certain cells that have been 5 genetically engineered, using methods such as those described herein, to express and secrete the polypeptide. Such cells may be animal or human cells, and may be autologous, heterologous, or xenogeneic. Optionally, the cells may be immortalized. In order to decrease the chance of an 10 immunological response, the cells may be encapsulated to avoid infiltration of surrounding tissues. The encapsulation materials are typically biocompatible, semi-permeable polymeric enclosures or membranes that allow the release of the protein product(s) but prevent the destruction of the 15 cells by the patient's immune system or by other detrimental factors from the surrounding tissues.

Additional embodiments of the present invention relate to cells and methods (e.g., homologous recombination and/or other recombinant production methods) for both the *in vitro* 20 production of therapeutic polypeptides and for the production and delivery of therapeutic polypeptides by gene therapy or cell therapy. Homologous and other recombination methods may be used to modify a cell that contains a normally transcriptionally silent IL-17 receptor like gene, or an under 25 expressed gene, and thereby produce a cell which expresses therapeutically efficacious amounts of IL-17 receptor like polypeptides.

It is further envisioned that IL-17 receptor like polypeptide can be produced *in vitro* or *in vivo* by homologous 30 recombination, or with recombinant production methods utilizing control elements introduced into cells already containing DNA encoding IL-17 receptor like polypeptides. For example, homologous recombination is a technique originally developed for targeting genes to induce or correct mutations

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in transcriptionally active genes (Kucherlapati, *Prog. in Nucl. Acid Res. & Mol. Biol.*, 36:301, 1989). The basic technique was developed as a method for introducing specific mutations into specific regions of the mammalian genome 5 (Thomas et al., *Cell*, 44:419-428, 1986; Thomas and Capecchi, *Cell*, 51:503-512, 1987; Doetschman et al., *Proc. Natl. Acad. Sci.*, 85:8583-8587, 1988) or to correct specific mutations within defective genes (Doetschman et al., *Nature*, 330:576-10 578, 1987). Exemplary homologous recombination techniques are described in U.S. Patent No. 5,272,071 (EP 9193051, EP Publication No. 505500; PCT/US90/07642, International Publication No. WO 91/09955).

Through homologous recombination, the DNA sequence to be inserted into the genome can be directed to a specific region 15 of the gene of interest by attaching it to targeting DNA. The targeting DNA is a nucleotide sequence that is complementary (homologous) to a region of the genomic DNA. Small pieces of targeting DNA that are complementary to a specific region of the genome are put in contact with the parental strand during 20 the DNA replication process. It is a general property of DNA that has been inserted into a cell to hybridize, and therefore, recombine with other pieces of endogenous DNA through shared homologous regions. If this complementary strand is attached to an oligonucleotide that contains a 25 mutation or a different sequence or an additional nucleotide, it too is incorporated into the newly synthesized strand as a result of the recombination. As a result of the proofreading function, it is possible for the new sequence of DNA to serve as the template. Thus, the transferred DNA is incorporated 30 into the genome.

Attached to these pieces of targeting DNA are regions of DNA which may interact with or control the expression of a IL-17 receptor like polypeptide, e.g., flanking sequences. For example, a promoter/enhancer element, a suppresser, or an

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exogenous transcription modulatory element is inserted in the genome of the intended host cell in proximity and orientation sufficient to influence the transcription of DNA encoding the desired IL-17 receptor like polypeptide. The control element
5 controls a portion of the DNA present in the host cell genome. Thus, the expression of the desired IL-17 receptor like polypeptide may be achieved not by transfection of DNA that encodes the IL-17 receptor like gene itself, but rather by the use of targeting DNA (containing regions of homology with the
10 endogenous gene of interest) coupled with DNA regulatory segments that provide the endogenous gene sequence with recognizable signals for transcription of an IL-17 receptor like polypeptide.

In an exemplary method, the expression of a desired
15 targeted gene in a cell (*i.e.*, a desired endogenous cellular gene) is altered via homologous recombination into the cellular genome at a preselected site, by the introduction of DNA which includes at least a regulatory sequence, an exon and a splice donor site. These components are introduced into the
20 chromosomal (genomic) DNA in such a manner that this, in effect, results in the production of a new transcription unit (in which the regulatory sequence, the exon and the splice donor site present in the DNA construct are operatively linked to the endogenous gene). As a result of the introduction of
25 these components into the chromosomal DNA, the expression of the desired endogenous gene is altered.

Altered gene expression, as described herein, encompasses activating (or causing to be expressed) a gene which is normally silent (unexpressed) in the cell as obtained, as well
30 as increasing the expression of a gene which is not expressed at physiologically significant levels in the cell as obtained. The embodiments further encompass changing the pattern of regulation or induction such that it is different from the pattern of regulation or induction that occurs in the cell as

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obtained, and reducing (including eliminating) the expression of a gene which is expressed in the cell as obtained.

One method by which homologous recombination can be used to increase, or cause, IL-17 receptor like polypeptide production from a cell's endogenous IL-17 receptor like gene involves first using homologous recombination to place a recombination sequence from a site-specific recombination system (e.g., Cre/loxP, FLP/FRT) (Sauer, *Current Opinion In Biotechnology*, 5:521-527, 1994; Sauer, *Methods In Enzymology*, 225:890-900, 1993) upstream (that is, 5' to) of the cell's endogenous genomic IL-17 receptor like polypeptide coding region. A plasmid containing a recombination site homologous to the site that was placed just upstream of the genomic IL-17 receptor like polypeptide coding region is introduced into the modified cell line along with the appropriate recombinase enzyme. This recombinase causes the plasmid to integrate, via the plasmid's recombination site, into the recombination site located just upstream of the genomic IL-17 receptor like polypeptide coding region in the cell line (Baubonis and Sauer, *Nucleic Acids Res.*, 21:2025-2029, 1993; O'Gorman et al., *Science*, 251:1351-1355, 1991). Any flanking sequences known to increase transcription (e.g., enhancer/promoter, intron, translational enhancer), if properly positioned in this plasmid, would integrate in such a manner as to create a new or modified transcriptional unit resulting in *de novo* or increased IL-17 receptor like polypeptide production from the cell's endogenous IL-17 receptor like gene.

A further method to use the cell line in which the site specific recombination sequence had been placed just upstream of the cell's endogenous genomic IL-17 receptor like polypeptide coding region is to use homologous recombination to introduce a second recombination site elsewhere in the cell line's genome. The appropriate recombinase enzyme is then introduced into the two-recombination-site cell line, causing

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a recombination event (deletion, inversion, translocation) (Sauer, *Current Opinion In Biotechnology*, *supra*, 1994; Sauer, *Methods In Enzymology*, *supra*, 1993) that would create a new or modified transcriptional unit resulting in *de novo* or 5 increased IL-17 receptor like polypeptide production from the cell's endogenous IL-17 receptor like gene.

An additional approach for increasing, or causing, the expression of IL-17 receptor like polypeptide from a cell's endogenous IL-17 receptor like gene involves increasing, or 10 causing, the expression of a gene or genes (e.g., transcription factors) and/or decreasing the expression of a gene or genes (e.g., transcriptional repressors) in a manner which results in *de novo* or increased IL-17 receptor like polypeptide production from the cell's endogenous IL-17 15 receptor like gene. This method includes the introduction of a non-naturally occurring polypeptide (e.g., a polypeptide comprising a site specific DNA binding domain fused to a transcriptional factor domain) into the cell such that *de novo* or increased IL-17 receptor like polypeptide production from 20 the cell's endogenous IL-17 receptor like gene results.

The present invention further relates to DNA constructs useful in the method of altering expression of a target gene. In certain embodiments, the exemplary DNA constructs comprise: (a) one or more targeting sequences; (b) a regulatory 25 sequence; (c) an exon; and (d) an unpaired splice-donor site. The targeting sequence in the DNA construct directs the integration of elements (a)-(d) into a target gene in a cell such that the elements (b)-(d) are operatively linked to sequences of the endogenous target gene. In another 30 embodiment, the DNA constructs comprise: (a) one or more targeting sequences, (b) a regulatory sequence, (c) an exon, (d) a splice-donor site, (e) an intron, and (f) a splice-acceptor site, wherein the targeting sequence directs the integration of elements (a)-(f) such that the elements of (b)-

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(f) are operatively linked to the endogenous gene. The targeting sequence is homologous to the preselected site in the cellular chromosomal DNA with which homologous recombination is to occur. In the construct, the exon is 5 generally 3' of the regulatory sequence and the splice-donor site is 3' of the exon.

If the sequence of a particular gene is known, such as the nucleic acid sequence of IL-17 receptor like polypeptide presented herein, a piece of DNA that is complementary to a 10 selected region of the gene can be synthesized or otherwise obtained, such as by appropriate restriction of the native DNA at specific recognition sites bounding the region of interest. This piece serves as a targeting sequence(s) upon insertion into the cell and will hybridize to its homologous region 15 within the genome. If this hybridization occurs during DNA replication, this piece of DNA, and any additional sequence attached thereto, will act as an Okazaki fragment and will be incorporated into the newly synthesized daughter strand of DNA. The present invention, therefore, includes nucleotides 20 encoding a IL-17 receptor like polypeptide, which nucleotides may be used as targeting sequences.

IL-17 receptor like polypeptide cell therapy, e.g., the implantation of cells producing IL-17 receptor like polypeptides, is also contemplated. This embodiment involves 25 implanting cells capable of synthesizing and secreting a biologically active form of IL-17 receptor like polypeptide. Such IL-17 receptor like polypeptide-producing cells can be cells that are natural producers of IL-17 receptor like polypeptides or may be recombinant cells whose ability to 30 produce IL-17 receptor like polypeptides has been augmented by transformation with a gene encoding the desired IL-17 receptor like polypeptide or with a gene augmenting the expression of IL-17 receptor like polypeptide. Such a modification may be accomplished by means of a vector suitable for delivering the

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gene as well as promoting its expression and secretion. In order to minimize a potential immunological reaction in patients being administered an IL-17 receptor like polypeptide, as may occur with the administration of a 5 polypeptide of a foreign species, it is preferred that the natural cells producing IL-17 receptor like polypeptide be of human origin and produce human IL-17 receptor like polypeptide. Likewise, it is preferred that the recombinant cells producing IL-17 receptor like polypeptide be transformed 10 with an expression vector containing a gene encoding a human IL-17 receptor like polypeptide.

Implanted cells may be encapsulated to avoid the infiltration of surrounding tissue. Human or non-human animal cells may be implanted in patients in biocompatible, 15 semipermeable polymeric enclosures or membranes that allow the release of IL-17 receptor like polypeptide, but that prevent the destruction of the cells by the patient's immune system or by other detrimental factors from the surrounding tissue. Alternatively, the patient's own cells, transformed to produce 20 IL-17 receptor like polypeptides *ex vivo*, may be implanted directly into the patient without such encapsulation.

Techniques for the encapsulation of living cells are known in the art, and the preparation of the encapsulated cells and their implantation in patients may be routinely 25 accomplished. For example, Baetge et al. (WO95/05452; PCT/US94/09299) describe membrane capsules containing genetically engineered cells for the effective delivery of biologically active molecules. The capsules are biocompatible and are easily retrievable. The capsules encapsulate cells 30 transfected with recombinant DNA molecules comprising DNA sequences coding for biologically active molecules operatively linked to promoters that are not subject to down regulation *in vivo* upon implantation into a mammalian host. The devices provide for the delivery of the molecules from living cells to

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specific sites within a recipient. In addition, see U.S. Patent Nos. 4,892,538, 5,011,472, and 5,106,627. A system for encapsulating living cells is described in PCT Application no. PCT/US91/00157 of Aebischer et al. See also, PCT Application no. PCT/US91/00155 of Aebischer et al., Winn et al., *Exper. Neurol.*, 113:322-329 (1991), Aebischer et al., *Exper. Neurol.*, 111:269-275 (1991); and Tresco et al., ASAIO, 38:17-23 (1992).

In vivo and in vitro gene therapy delivery of IL-17 receptor like polypeptides is also envisioned. One example of a gene therapy technique is to use the IL-17 receptor like gene (either genomic DNA, cDNA, and/or synthetic DNA) encoding a IL-17 receptor like polypeptide which may be operably linked to a constitutive or inducible promoter to form a "gene therapy DNA construct". The promoter may be homologous or heterologous to the endogenous IL-17 receptor like gene, provided that it is active in the cell or tissue type into which the construct will be inserted. Other components of the gene therapy DNA construct may optionally include, DNA molecules designed for site-specific integration (e.g., endogenous sequences useful for homologous recombination), tissue-specific promoter, enhancer(s) or silencer(s), DNA molecules capable of providing a selective advantage over the parent cell, DNA molecules useful as labels to identify transformed cells, negative selection systems, cell specific binding agents (as, for example, for cell targeting), cell-specific internalization factors, and transcription factors to enhance expression by a vector as well as factors to enable vector manufacture.

A gene therapy DNA construct can then be introduced into cells (either ex vivo or in vivo) using viral or non-viral vectors. One means for introducing the gene therapy DNA construct is by means of viral vectors as described herein. Certain vectors, such as retroviral vectors, will deliver the DNA construct to the chromosomal DNA of the cells, and the

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gene can integrate into the chromosomal DNA. Other vectors will function as episomes, and the gene therapy DNA construct will remain in the cytoplasm:

In yet other embodiments, regulatory elements can be
5 included for the controlled expression of the IL-17 receptor
like gene in the target cell. Such elements are turned on in
response to an appropriate effector. In this way, a
therapeutic polypeptide can be expressed when desired. One
conventional control means involves the use of small molecule
10 dimerizers or rapalogs (as described in WO9641865
(PCT/US96/099486); WO9731898 (PCT/US97/03137) and WO9731899
(PCT/US95/03157) used to dimerize chimeric proteins which
contain a small molecule-binding domain and a domain capable
of initiating biological process, such as a DNA-binding
15 protein or transcriptional activation protein. The
dimerization of the proteins can be used to initiate
transcription of the transgene.

An alternative regulation technology uses a method of
storing proteins expressed from the gene of interest inside
20 the cell as an aggregate or cluster. The gene of interest is
expressed as a fusion protein that includes a conditional
aggregation domain which results in the retention of the
aggregated protein in the endoplasmic reticulum. The stored
proteins are stable and inactive inside the cell. The
25 proteins can be released, however, by administering a drug
(e.g., small molecule ligand) that removes the conditional
aggregation domain and thereby specifically breaks apart the
aggregates or clusters so that the proteins may be secreted
from the cell. See, *Science* 287:816-817, and 826-830 (2000).

30 Other suitable control means or gene switches include,
but are not limited to, the following systems. Mifepristone
(RU486) is used as a progesterone antagonist. The binding of
a modified progesterone receptor ligand-binding domain to the

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progesterone antagonist activates transcription by forming a dimer of two transcription factors which then pass into the nucleus to bind DNA. The ligand binding domain is modified to eliminate the ability of the receptor to bind to the natural 5 ligand. The modified steroid hormone receptor system is further described in U.S. 5,364,791; WO9640911, and WO9710337.

Yet another control system uses ecdysone (a fruit fly steroid hormone) which binds to and activates an ecdysone receptor (cytoplasmic receptor). The receptor then 10 translocates to the nucleus to bind a specific DNA response element (promoter from ecdysone-responsive gene). The ecdysone receptor includes a transactivation domain/DNA-binding domain/ligand-binding domain to initiate transcription. The ecdysone system is further described in. 15 U.S. 5,514,578; WO9738117; WO9637609; and WO9303162.

Another control means uses a positive tetracycline-controllable transactivator. This system involves a mutated tet repressor protein DNA-binding domain (mutated tet R-4 amino acid changes which resulted in a reverse tetracycline- 20 regulated transactivator protein, i.e., it binds to a tet operator in the presence of tetracycline) linked to a polypeptide which activates transcription. Such systems are described in U.S. Patent Nos. 5,464,758; 5,650,298 and 5,654,168.

25 Additional expression control systems and nucleic acid constructs are described in U.S. Patent Nos. 5,741,679 and 5,834,186, to Innovir Laboratories Inc.

In vivo gene therapy may be accomplished by introducing the gene encoding an IL-17 receptor like polypeptide into 30 cells via local injection of an IL-17 receptor like nucleic acid molecule or by other appropriate viral or non-viral delivery vectors. Hefti, *Neurobiology*, 25:1418-1435 (1994). For example, a nucleic acid molecule encoding an IL-17

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receptor like polypeptide may be contained in an adeno-associated virus (AAV) vector for delivery to the targeted cells (e.g., Johnson, International Publication No. WO95/34670; International Application No. PCT/US95/07178).

5 The recombinant AAV genome typically contains AAV inverted terminal repeats flanking a DNA sequence encoding an IL-17 receptor like polypeptide operably linked to functional promoter and polyadenylation sequences.

Alternative suitable viral vectors include, but are not limited to, retrovirus, adenovirus, herpes simplex virus, lentivirus, hepatitis virus, parvovirus, papovavirus, poxvirus, alphavirus, coronavirus, rhabdovirus, paramyxovirus, and papilloma virus vectors. U.S. Patent No. 5,672,344 describes an *in vivo* viral-mediated gene transfer system involving a recombinant neurotrophic HSV-1 vector. U.S. Patent No. 5,399,346 provides examples of a process for providing a patient with a therapeutic protein by the delivery of human cells which have been treated *in vitro* to insert a DNA segment encoding a therapeutic protein. Additional methods and materials for the practice of gene therapy techniques are described in U.S. Patent No. 5,631,236 involving adenoviral vectors; U.S. Patent No. 5,672,510 involving retroviral vectors; and U.S. 5,635,399 involving retroviral vectors expressing cytokines.

25 Nonviral delivery methods include, but are not limited to, liposome-mediated transfer, naked DNA delivery (direct injection), receptor-mediated transfer (ligand-DNA complex), electroporation, calcium phosphate precipitation, and microparticle bombardment (e.g., gene gun). Gene therapy materials and methods may also include the use of inducible promoters, tissue-specific enhancer-promoters, DNA sequences designed for site-specific integration, DNA sequences capable of providing a selective advantage over the parent cell, labels to identify transformed cells, negative selection

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systems and expression control systems (safety measures), cell-specific binding agents (for cell targeting), cell-specific internalization factors, and transcription factors to enhance expression by a vector as well as methods of vector manufacture. Such additional methods and materials for the practice of gene therapy techniques are described in U.S. Patent No. 4,970,154 involving electroporation techniques; WO96/40958 involving nuclear ligands; U.S. Patent No. 5,679,559 describing a lipoprotein-containing system for gene delivery; U.S. Patent No. 5,676,954 involving liposome carriers; U.S. Patent No. 5,593,875 concerning methods for calcium phosphate transfection; and U.S. Patent No. 4,945,050 wherein biologically active particles are propelled at cells at a speed whereby the particles penetrate the surface of the cells and become incorporated into the interior of the cells.

It is also contemplated that IL-17 receptor like gene therapy or cell therapy can further include the delivery of one or more additional polypeptide(s) in the same or a different cell(s). Such cells may be separately introduced into the patient, or the cells may be contained in a single implantable device, such as the encapsulating membrane described above, or the cells may be separately modified by means of viral vectors.

A means to increase endogenous IL-17 receptor like polypeptide expression in a cell via gene therapy is to insert one or more enhancer elements into the IL-17 receptor like polypeptide promoter, where the enhancer element(s) can serve to increase transcriptional activity of the IL-17 receptor like gene. The enhancer element(s) used will be selected based on the tissue in which one desires to activate the gene(s); enhancer elements known to confer promoter activation in that tissue will be selected. For example, if a gene encoding a IL-17 receptor like polypeptide is to be "turned on" in T-cells, the *lck* promoter enhancer element may be used.

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Here, the functional portion of the transcriptional element to be added may be inserted into a fragment of DNA containing the IL-17 receptor like polypeptide promoter (and optionally, inserted into a vector and/or 5' and/or 3' flanking sequence(s), etc.) using standard cloning techniques. This construct, known as a "homologous recombination construct", can then be introduced into the desired cells either ex vivo or in vivo.

Gene therapy also can be used to decrease IL-17 receptor like polypeptide expression by modifying the nucleotide sequence of the endogenous promoter(s). Such modification is typically accomplished via homologous recombination methods. For example, a DNA molecule containing all or a portion of the promoter of the IL-17 receptor like gene(s) selected for inactivation can be engineered to remove and/or replace pieces of the promoter that regulate transcription. For example the TATA box and/or the binding site of a transcriptional activator of the promoter may be deleted using standard molecular biology techniques; such deletion can inhibit promoter activity thereby repressing the transcription of the corresponding IL-17 receptor like gene. The deletion of the TATA box or the transcription activator binding site in the promoter may be accomplished by generating a DNA construct comprising all or the relevant portion of the IL-17 receptor like polypeptide promoter(s) (from the same or a related species as the IL-17 receptor like gene(s) to be regulated) in which one or more of the TATA box and/or transcriptional activator binding site nucleotides are mutated via substitution, deletion and/or insertion of one or more nucleotides. As a result, the TATA box and/or activator binding site has decreased activity or is rendered completely inactive. The construct will typically contain at least about 500 bases of DNA that correspond to the native (endogenous) 5' and 3' DNA sequences adjacent to the promoter segment that has been modified. The construct may be introduced into the

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appropriate cells (either *ex vivo* or *in vivo*) either directly or via a viral vector as described herein. Typically, the integration of the construct into the genomic DNA of the cells will be via homologous recombination, where the 5' and 3' DNA sequences in the promoter construct can serve to help integrate the modified promoter region via hybridization to the endogenous chromosomal DNA.

10 Additional Uses of IL-17 receptor like Nucleic Acids and Polypeptides

Nucleic acid molecules of the present invention (including those that do not themselves encode biologically active polypeptides) may be used to map the locations of the IL-17 receptor like gene and related genes on chromosomes. 15 Mapping may be done by techniques known in the art, such as PCR amplification and *in situ* hybridization:

IL-17 receptor like nucleic acid molecules (including those that do not themselves encode biologically active polypeptides), may be useful as hybridization probes in 20 diagnostic assays to test, either qualitatively or quantitatively, for the presence of an IL-17 receptor like DNA or corresponding RNA in mammalian tissue or bodily fluid samples.

The IL-17 receptor like polypeptides may be used 25 (simultaneously or sequentially) in combination with one or more cytokines, growth factors, antibiotics, anti-inflammatories, and/or chemotherapeutic agents as is appropriate for the indication being treated.

Other methods may also be employed where it is desirable 30 to inhibit the activity of one or more IL-17 receptor like polypeptides. Such inhibition may be effected by nucleic acid molecules which are complementary to and hybridize to expression control sequences (triple helix formation) or to IL-17 receptor like mRNA. For example, antisense DNA or RNA

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molecules, which have a sequence that is complementary to at least a portion of the selected IL-17 receptor like gene(s) can be introduced into the cell. Antisense probes may be designed by available techniques using the sequence of IL-17 receptor like polypeptide disclosed herein. Typically, each such antisense molecule will be complementary to the start site (5' end) of each selected IL-17 receptor like gene. When the antisense molecule then hybridizes to the corresponding IL-17 receptor like mRNA, translation of this mRNA is prevented or reduced. Antisense inhibitors provide information relating to the decrease or absence of an IL-17 receptor like polypeptide in a cell or organism.

Alternatively, gene therapy may be employed to create a dominant-negative inhibitor of one or more IL-17 receptor like polypeptides. In this situation, the DNA encoding a mutant polypeptide of each selected IL-17 receptor like polypeptide can be prepared and introduced into the cells of a patient using either viral or non-viral methods as described herein. Each such mutant is typically designed to compete with endogenous polypeptide in its biological role.

In addition, an IL-17 receptor like polypeptide, whether biologically active or not, may be used as an immunogen, that is, the polypeptide contains at least one epitope to which antibodies may be raised. Selective binding agents that bind to an IL-17 receptor like polypeptide (as described herein) may be used for *in vivo* and *in vitro* diagnostic purposes, including, but not limited to, use in labeled form to detect the presence of IL-17 receptor like polypeptide in a body fluid or cell sample. The antibodies may also be used to prevent, treat, or diagnose a number of diseases and disorders, including those recited herein. The antibodies may bind to an IL-17 receptor like polypeptide so as to diminish or block at least one activity characteristic of an IL-17 receptor like polypeptide, or may bind to a polypeptide to increase at least one activity characteristic of an IL-17

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receptor like polypeptide (including by increasing the pharmacokinetics of the IL-17 receptor like polypeptide).

EXAMPLE 1

5 Cloning of cDNA Encoding human IL-17 Receptor Like Polypeptide

Materials and methods for cDNA cloning and analysis are described in Sambrook et al., supra. which is incorporated herein by reference..

An EST fragment zhgb-a1133097, was identified in an
10 internal database. The fragment encoded a partial peptide sequence that is related to the human and mouse IL-17 receptor. The EST sequence was used to generate two gene specific primers, 2418-63 (5' CGA GCC ATG CTG GCT GAT GTT C
15 3': SEQ ID NO: 8) and 2418-64 (5' CGT GGT CGA AGG ACA CCT GCA TG 3'; SEQ ID NO: 6), which correspond to the 5' region of the fragment. These primers were used to extend the 5' region of the fragment.

To isolate the full length cDNA, PCR was used to screen a
20 panel of 77 human tissue libraries. The PCR was carried out with Ready-To-Go Beads (Amersham Pharmacia Biotech, cat no. 27-9553-01), 50 ng of cDNA library and 5 pmol of each of the following primers: 2714-51 (5' CCA GTG TTT CGC CTA CTT CCT CC
25 3'; SEQ ID NO: 4) and 2417-56 (5' GAT ATC CGG TAA AGG GTT GGG GC 3'; SEQ ID NO: 5). The PCR reactions were carried out at 94°C for 30 seconds, 5 cycles of 94°C for 15 seconds, 72°C for 2 minutes, then 5 cycles of 94°C for 15 seconds, 70°C for 2 minutes, followed by 25 cycles at 94°C for 15 seconds, 68°C for 2.5 minutes. Positive cDNA libraries expressed a 1247 bp
30 band. The following ten cDNA libraries were scored as positive: fetal pancreas random library, fetal kidney oligo-dT library, fetal ovary oligo-dT libarary, fetal calveria oligo-dT library, fetal femur oligo-dT library, fetal gall bladder oligo-dT library, fetal gall bladder random library, spinal column oligo-dT library, spinal column random
35 library,

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library, and bone (limb) oligo-dT library.

To further characterize the 10 positive libraries, 5'RACE PCR was carried out with 25 ng of cDNA from each library, 200 μ M dNTP , 1x Advantage cDNA Polymerase Mix (Clontech, CA cat.

5 No. 8417-1) and 10 pmol of a gene specific primer 2418-64 (5' CGT GGT CGA AGG ACA CCT GCA TG 3'; SEQ ID NO: 6) and 10 pmol of a pSPORT vector primer 1916-83 (5' GGC TGT ATG TTG TGT GGA ATT GTG AGC G 3'; SEQ ID NO: 7) in a final volume of 50 μ l. The PCR reactions were carried out as follows: 94°C for 2

10 minutes, 5 cycles of 94°C for 15 seconds and 72°C for 4 minutes, then 5 cycles of 94°C for 15 seconds and 70°C for 4 minutes followed by 25 cycles of 94°C for 15 seconds and 68°C for 4 minutes.

The resulting PCR products were re-amplified using nested primers. Briefly, 5 μ l of 1:50 dilution of each product was used along with 10 pmol of the gene specific primer 2418-63 (5' CGA GCC ATG CTG GCT GAT GTT C 3'; SEQ ID NO: 8) and vector nest primer 1916-82 (5' CAT GAT TA CGC CAA GCT CTA ATA CGA CTC 3'; SEQ ID NO: 9) in each reaction. The RACE reactions were 20 carried out as described above for the primary RACE reaction. The final RACE products (8 μ l) was analyzed on 1% TAE agarose gel at 5V/cm. The well-defined single bands were purified using the QIAquick PCR purification kit (QIAGen, Cat. No. 28104) and sequenced using standard methods.

25 For the second round of 5' RACE, another gene specific primer 2432-38 (5' GAA GCT ACT GTT GAG CTG CTT CG 3'; SEQ ID NO: 10) and the pCMV/SPORT vector primer 2182-36 (5' CCG ATC CAG CCT CCG GAC TCT AG 3'; SEQ ID NO: 11) were used. Plasmid DNAs of LTI human multi-tissue cDNA library were used as 30 templates. The PCR reactions were carried out in 50 μ l volume containing 30 ng of cDNA for each library pool, 10 pmol of each primer, 200 μ M dNTP and a 1x Advantage cDNA Polymerase Mix (Clontech, cat. No. 8417-1). The PCR reactions were carried out as follows: 94°C for 1 minute, followed by 35

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cycles of 94°C for 15 seconds, 65°C for 2 minutes and 72°C for 1 minute, followed by a final extension of 72°C for 10 minutes.

The resulting products were again re-amplified using 5 another pair of nested primers, a gene specific primer 2144-06 (5' GCGTCAGCAATCACATGCTTCCC 3'; SEQ ID NO: 12) and the vector primer 2144-06 (5' GCCTATTAGGTGACACTATAGAAC 3'; SEQ ID NO: 13). The PCR reactions were carried out as described above. The final products were analyzed by agarose gel 10 electrophoresis, subcloned into the PCR 2.1-TOPO vector (Invitrogen Cat. No. K4560-01) and the inserts were sequences using standard methods.

The resulting cDNA is set out as SEQ ID NO: 1 and is 3083 nucleotides in length. This cDNA contains a 2214 nucleotide 15 coding region and encodes a 738 amino acid polypeptide which is set out as SEQ ID NO: 2 and is denoted as IL-17 receptor like polypeptide.

EXAMPLE 2

20 Tissue Expression of IL-17 Receptor like Polynucleotide

Quantitative PCR was carried out on various human fetal tissues to analyze the expression pattern of IL-17 receptor like mRNA. Total RNA was isolated with the Total RNA Isolation kit (Amersham Pharmacia Biotech, cat. No. 15593-031). The reverse transcriptase reactions were carried out as follows: 2 µg of total RNA was mixed with 1 µl (50 ng/µl) of random primer and incubated at 70°C for 10 minutes then quick-chilled on ice. Subsequently, 4 µl of 5x First Stand Buffer (BRL), 2 µl of 0.1 M DTT (BRL) and 1 µl of dNTP mix was added 25 to the reactions and mixed well. The reactions were incubated at 37°C for 2 minutes. Subsequently, 1 µl of Superscript II reverse transcriptase (BRL) was added and incubated at 37°C, for 1 hour. The reactions were terminated by placing the tubes on ice.

30 The subsequent PCR reactions were carried using Ready-To-

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Go PCR Bead plates (Amersham Pharmacia Biotech cat. No. 27-9553-01). The following components were added to each well: 1 µl of the RT reaction mixture after normalization with G#PDH levels, 1 µl (10 pmol/µl) of primer 2417-51 (5' CCA GTG TTT CGC CTA CCT CCT CC 3'; SEQ ID NO: 14), 1 µl (10 pmol/µl) of primer 2418-65 (5' GGA GCT TTT CGG CAA TGG CTG AC 3'; SEQ ID NO: 15) and 22 µl of water. The reactions were mixed well carried out as follows: 94°C for 1 minute, followed by 5 cycles of 94°C for 30 seconds, 72°C for 4 minutes, 5 cycles of 94°C for 30 seconds, 70°C for 4 minutes and 25 cycles of 94°C for 30 seconds and 68°C for 4 minutes.

Expression of IL-17 receptor like transcript was detected on agarose gels and the intensities of expression as listed below. "++++" indicates a strong signal while "+" denotes a weak signal and "-" indicates no expression.

SOURCE	SIGNAL
Testis	++++
Kidney	++++
Intestine	+++
Pancreas	+++
Spinal Cord	+++
Bone	+++
Thymus	+
Placenta	-

20

EXAMPLE 3

Production of IL-17 Receptor like Polypeptides

A. Bacterial Expression of IL-17 Receptor like Polypeptides

PCR is used to amplify template DNA sequences encoding a IL-17 receptor like polypeptide using primers corresponding to

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the 5' and 3' ends of the sequence. The amplified DNA products may be modified to contain restriction enzyme sites to allow for insertion into expression vectors. PCR products are gel purified and inserted into expression vectors using standard recombinant DNA methodology. An exemplary vector, such as 5 pAMG21 (ATCC No. 98113) containing the lux promoter and a gene encoding kanamycin resistance is digested with BamHI and NdeI for directional cloning of inserted DNA. The ligated mixture is transformed into an *E. coli* host strain by electroporation 10 and transformants are selected for kanamycin resistance. Plasmid DNA from selected colonies is isolated and subjected to DNA sequencing to confirm the presence of the insert.

Transformed host cells are incubated in 2x YT medium containing 30 mg/ml kanamycin at 30°C prior to induction. 15 Gene expression is induced by the addition of N-(3-oxohexanoyl)-dl-homoserine lactone to a final concentration of 30 ng/ml followed by incubation at either 30°C or 37°C for six hours. The expression of IL-17 receptor like polypeptide is evaluated by centrifugation of the culture, resuspension and 20 lysis of the bacterial pellets, and analysis of host cell proteins by SDS-polyacrylamide gel electrophoresis.

Inclusion bodies containing IL-17 receptor like polypeptide are purified as follows. Bacterial cells are pelleted by centrifugation and resuspended in water. The cell 25 suspension is lysed by sonication and pelleted by centrifugation at 195,000 x g for 5 to 10 minutes. The supernatant is discarded, and the pellet is washed and transferred to a homogenizer. The pellet is homogenized in 5 ml of a Percoll solution (75% liquid Percoll/0.15 M NaCl) 30 until uniformly suspended and then diluted and centrifuged at 21,600 x g for 30 minutes. Gradient fractions containing the inclusion bodies are recovered and pooled. The isolated inclusion bodies are analyzed by SDS-PAGE. A single band on an SDS polyacrylamide gel corresponding to *E. coli*-produced 35 IL-17 receptor like polypeptide is excised from the gel, and

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the N-terminal amino acid sequence is determined essentially as described by Matsudaira et al., J. Biol. Chem., 262: 10-35 (1987).

5 B. Mammalian Cell Production of IL-17 Receptor like Polypeptides

The IL-17 receptor like DNA was subcloned into a mammalian expression vector as described above using standard DNA technology. An exemplary expression vector, pCEP4 (Invitrogen, Carlsbad, CA), which contains an Epstein-Barr virus origin of replication, may be used for the expression of IL-17 receptor like in 293-EBNA-1 cells. Amplified and gel purified PCR products are ligated into pCEP4 vector and lipofected into 293-EBNA cells. The transfected cells are selected in 100 mg/ml hygromycin and the resulting drug-resistant cultures are grown to confluence. The cells are then cultured in serum-free media for 72 hours. The conditioned media is removed and, IL-17 receptor like protein polypeptide expression is analyzed by SDS-PAGE. Human E3 α polypeptide expression may be detected by silver staining. Alternatively, IL-17 receptor like polypeptide is produced as a fusion protein with an epitope tag, such as an IgG constant domain or a FLAG epitope, which may be detected by Western blot analysis using antibodies to the tag peptide.

25 Human E3 α polypeptides may be excised from an SDS-polyacrylamide gel, or IL-17 receptor like fusion proteins are purified by affinity chromatography to the epitope tag, and subjected to N-terminal amino acid sequence analysis as described herein.

30

EXAMPLE 4

Production of Anti- IL-17 receptor like Polypeptide

Antibodies

35 Polyclonal or monoclonal antibodies to IL-17 receptor like polypeptides may be obtained by immunization of animals

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with purified protein or with IL-17 receptor like peptides produced by biological or chemical synthesis. Suitable procedures for generating antibodies include those described in Hudson and Bay, *Practical Immunology, Second Edition*,

5 Edition, Blackwell Scientific Publications.

In one procedure for the production of monoclonal antibodies, animals (typically mice or rabbits) are injected with a IL-17 receptor like polypeptide antigens (such as an recombinant full length or truncated forms of IL-17 receptor 10 like polypeptide, analogs, variants or the like), and those with sufficient serum titer levels as determined by ELISA are selected for hybridoma production. Spleens of immunized animals are collected and prepared as single cell suspensions from which splenocytes are recovered. The splenocytes are 15 fused to mouse myeloma cells (such as Sp2/0-Ag14 cells), allowed to incubate in DMEM with 200 U/ml penicillin, 200 mg/ml streptomycin sulfate, and 4 mM glutamine, then incubated in HAT selection medium (Hypoxanthine; Aminopterin; Thymidine). After selection, the tissue culture supernatants 20 are taken from each

fusion well and tested for anti- IL-17 receptor like polypeptide antibody production by ELISA.

Alternative procedures for obtaining anti- IL-17 receptor 25 like polypeptide antibodies may also be employed, such as the immunization of transgenic mice harboring human Ig loci for production of human antibodies, and the screening of synthetic antibody libraries, such as those generated by mutagenesis of an antibody variable domain.

-1-

SEQUENCE LISTING

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-2-

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Pro	Phe	Phe	Phe	Arg	Thr	Arg	Ala	Cys	Asp	Leu	Leu	Gln	Pro	Asp	
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Asn	Leu	Ala	Cys	Lys	Pro	Phe	Trp	Lys	Pro	Arg	Asn	Leu	Asn	Ile	Ser
	195					200			205						
Gln	His	Gly	Ser	Asp	Met	Gln	Val	Ser	Phe	Asp	His	Ala	Pro	His	Asn
	210					215			220						
Phe	Gly	Phe	Arg	Phe	Phe	Tyr	Leu	His	Tyr	Lys	Leu	Lys	His	Glu	Gly
	225					230			235			240			
Pro	Phe	Lys	Arg	Lys	Thr	Cys	Lys	Gln	Glu	Gln	Thr	Thr	Glu	Thr	Thr
	245					250			255						
Ser	Cys	Leu	Leu	Gln	Asn	Val	Ser	Pro	Gly	Asp	Tyr	Ile	Ile	Glu	Leu
	260					265			270						
Val	Asp	Asp	Thr	Asn	Thr	Thr	Arg	Lys	Val	Met	His	Tyr	Ala	Leu	Lys
	275					280			285						
Pro	Val	His	Ser	Pro	Trp	Ala	Gly	Pro	Ile	Arg	Ala	Met	Ala	Ile	Thr
	290					295			300						
Val	Pro	Leu	Val	Val	Ile	Ser	Ala	Phe	Ala	Thr	Leu	Phe	Thr	Val	Met
	305					310			315			320			

Cys Arg Lys Gln Gln Glu Asn Ile Tyr Ser His Leu Asp Glu Glu
 325 330 335
 Ser Ser Glu Ser Ser Thr Tyr Thr Ala Ala Leu Pro Arg Glu Arg Leu
 340 345 350
 Arg Pro Arg Pro Lys Val Phe Leu Cys Tyr Ser Ser Lys Asp Gly Gln
 355 360 365
 Asn His Met Asn Val Val Gln Cys Phe Ala Tyr Phe Leu Gln Asp Phe
 370 375 380
 Cys Gly Cys Glu Val Ala Leu Asp Leu Trp Glu Asp Phe Ser Leu Cys
 385 390 395 400
 Arg Glu Gly Gln Arg Glu Trp Val Ile Gln Lys Ile His Glu Ser Gln
 405 410 415
 Phe Ile Ile Val Val Cys Ser Lys Gly Met Lys Tyr Phe Val Asp Lys
 420 425 430
 Lys Asn Tyr Lys His Lys Gly Gly Arg Gly Ser Gly Lys Gly Glu
 435 440 445
 Leu Phe Leu Val Ala Val Ser Ala Ile Ala Glu Lys Leu Arg Gln Ala
 450 455 460
 Lys Gln Ser Ser Ser Ala Ala Leu Ser Lys Phe Ile Ala Val Tyr Phe
 465 470 475 480
 Asp Tyr Ser Cys Glu Gly Asp Val Pro Gly Ile Leu Asp Leu Ser Thr
 485 490 495
 Lys Tyr Arg Leu Met Asp Asn Leu Pro Gln Leu Cys Ser His Leu His
 500 505 510
 Ser Arg Asp His Gly Leu Gln Glu Pro Gly Gln His Thr Arg Gln Gly
 515 520 525
 Ser Arg Arg Asn Tyr Phe Arg Ser Lys Ser Gly Arg Ser Leu Tyr Val
 530 535 540
 Ala Ile Cys Asn Met His Gln Phe Ile Asp Glu Glu Pro Asp Trp Phe
 545 550 555 560
 Glu Lys Gln Phe Val Pro Phe His Pro Pro Pro Leu Arg Tyr Arg Glu
 565 570 575
 Pro Val Leu Glu Lys Phe Asp Ser Gly Leu Val Leu Asn Asp Val Met
 580 585 590
 Cys Lys Pro Gly Pro Glu Ser Asp Phe Cys Leu Lys Val Glu Ala Pro
 595 600 605
 Val Leu Gly Ala Thr Gly Pro Ala Asp Ser Gln His Glu Ser Gln His
 610 615 620
 Gly Gly Leu Asp Gln Asp Gly Glu Ala Arg Pro Ala Leu Asp Gly Ser
 625 630 635 640
 Ala Ala Leu Gln Pro Leu Leu His Thr Val Lys Ala Gly Ser Pro Ser
 645 650 655

Asp Met Pro Arg Asp Ser Gly Ile Tyr Asp Ser Ser Val Pro Ser Ser
 660 665 670

Glu Leu Ser Leu Pro Leu Met Glu Gly Leu Ser Thr Asp Gln Thr Glu
 675 680 685

Thr Ser Ser Leu Thr Glu Ser Val Ser Ser Ser Gly Leu Gly Glu
 690 695 700

Glu Glu Pro Pro Ala Leu Pro Ser Lys Leu Leu Ser Ser Gly Ser Cys
 705 710 715 720

Lys Ala Asp Leu Gly Cys Arg Ser Tyr Thr Asp Glu Leu His Ala Val
 725 730 735

Ala Pro

<210> 3

<211> 810

<212> PRT

<213> Homo sapiens

<400> 3

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Gly Leu Leu Leu Leu Leu Gly Val Leu Ala Pro Gly Gly Ala Ser
 20 25 30

Leu Arg Leu Leu Asp His Arg Ala Leu Val Cys Ser Gln Pro Gly Leu
 35 40 45

Asn Cys Thr Val Lys Asn Ser Thr Cys Leu Asp Asp Ser Trp Ile His
 50 55 60

Pro Arg Asn Leu Thr Pro Ser Ser Pro Lys Asp Leu Gln Ile Gln Leu
 65 70 75 80

His Phe Ala His Thr Gln Gln Gly Asp Leu Phe Pro Val Ala His Ile
 85 90 95

Glu Trp Thr Leu Gln Thr Asp Ala Ser Ile Leu Tyr Leu Glu Gly Ala
 100 105 110

Glu Leu Ser Val Leu Gln Leu Asn Thr Asn Glu Arg Leu Cys Val Arg
 115 120 125

Phe Glu Phe Leu Ser Lys Leu Arg His His Arg Arg Trp Arg Phe
 130 135 140

Thr Phe Ser His Phe Val Val Asp Pro Asp Gln Glu Tyr Glu Val Thr
 145 150 155 160

Val His His Leu Pro Lys Pro Ile Pro Asp Gly Asp Pro Asn His Gln
 165 170 175

Ser Lys Asn Phe Leu Val Pro Asp Cys Glu His Ala Arg Met Lys Val
 180 185 190

Thr Thr Pro Cys Met Ser Ser Gly Ser Leu Trp Asp Pro Asn Ile Thr
 195 200 205

Val Glu Thr Leu Glu Ala His Gln Leu Arg Val Ser Phe Thr Leu Trp
 210 215 220
 Asn Glu Ser Thr His Tyr Gln Ile Leu Leu Thr Ser Phe Pro His Met
 225 230 235 240
 Glu Asn His Ser Cys Phe Glu His Met His His Ile Pro Ala Pro Arg
 245 250 255
 Pro Glu Glu Phe His Gln Arg Ser Asn Val Thr Leu Thr Leu Arg Asn
 260 265 270
 Leu Lys Gly Cys Cys Arg His Gln Val Gln Ile Gln Pro Phe Phe Ser
 275 280 285
 Ser Cys Leu Asn Asp Cys Leu Arg His Ser Ala Thr Val Ser Cys Pro
 290 295 300
 Glu Met Pro Asp Thr Pro Glu Pro Ile Pro Asp Tyr Met Pro Leu Trp
 305 310 315 320
 Val Tyr Trp Phe Ile Thr Gly Ile Ser Ile Leu Leu Val Gly Ser Val
 325 330 335
 Ile Leu Leu Ile Val Cys Met Thr Trp Arg Leu Ala Gly Pro Gly Ser
 340 345 350
 Glu Lys Tyr Ser Asp Asp Thr Lys Tyr Thr Asp Gly Leu Pro Ala Ala
 355 360 365
 Asp Leu Ile Pro Pro Leu Lys Pro Arg Lys Val Trp Ile Ile Tyr
 370 375 380
 Ser Ala Asp His Pro Leu Tyr Val Asp Val Val Leu Lys Phe Ala Gln
 385 390 395 400
 Phe Leu Leu Thr Ala Cys Gly Thr Glu Val Ala Leu Asp Leu Leu Glu
 405 410 415
 Glu Gln Ala Ile Ser Glu Ala Gly Val Met Thr Trp Val Gly Arg Gln
 420 425 430
 Lys Gln Glu Met Val Glu Ser Asn Ser Lys Ile Ile Val Leu Cys Ser
 435 440 445
 Arg Gly Thr Arg Ala Lys Trp Gln Ala Leu Leu Gly Arg Gly Ala Pro
 450 455 460
 Val Arg Leu Arg Cys Asp His Gly Lys Pro Val Gly Asp Leu Phe Thr
 465 470 475 480
 Ala Ala Met Asn Met Ile Leu Pro Asp Phe Lys Arg Pro Ala Cys Phe
 485 490 495
 Gly Thr Tyr Val Val Cys Tyr Phe Ser Glu Val Ser Cys Asp Gly Asp
 500 505 510
 Val Pro Asp Leu Phe Gly Ala Ala Pro Arg Tyr Pro Leu Met Asp Arg
 515 520 525
 Phe Glu Glu Val Tyr Phe Arg Ile Gln Asp Leu Glu Met Phe Gln Pro
 530 535 540

Gly Arg Met His Arg Val Gly Glu Leu Ser Gly Asp Asn Tyr Leu Arg
 545 550 555 560

Ser Pro Gly Gly Arg Gln Leu Arg Ala Ala Leu Asp Arg Phe Arg Asp
 565 570 575

Trp Gln Val Arg Cys Pro Asp Trp Phe Glu Cys Glu Asn Leu Tyr Ser
 580 585 590

Ala Asp Asp Gln Asp Ala Pro Ser Leu Asp Glu Glu Val Phe Glu Glu
 595 600 605

Pro Leu Leu Pro Pro Gly Thr Gly Ile Val Lys Arg Ala Pro Leu Val
 610 615 620

Arg Glu Pro Gly Ser Gln Ala Cys Leu Ala Ile Asp Pro Leu Val Gly
 625 630 635 640

Glu Glu Gly Ala Ala Val Ala Lys Leu Glu Pro His Leu Gln Pro
 645 650 655

Arg Gly Gln Pro Ala Pro Gln Pro Leu His Thr Leu Val Leu Ala Ala
 660 665 670

Glu Glu Gly Ala Leu Val Ala Ala Val Glu Pro Gly Pro Leu Ala Asp
 675 680 685

Gly Ala Ala Val Arg Leu Ala Leu Ala Gly Glu Gly Glu Ala Cys Pro
 690 695 700

Leu Leu Gly Ser Pro Gly Ala Gly Arg Asn Ser Val Leu Phe Leu Pro
 705 710 715 720

Val Asp Pro Glu Asp Ser Pro Leu Gly Ser Ser Thr Pro Met Ala Ser
 725 730 735

Pro Asp Leu Leu Pro Glu Asp Val Arg Glu His Leu Glu Gly Leu Met
 740 745 750

Leu Ser Leu Phe Glu Gln Ser Leu Ser Cys Gln Ala Gln Gly Gly Cys
 755 760 765

Ser Arg Pro Ala Met Val Leu Thr Asp Pro His Thr Pro Tyr Glu Glu
 770 775 780 785

Glu Gln Arg Gln Ser Val Gln Ser Asp Gln Gly Tyr Ile Ser Arg Ser
 785 790 795 800

Ser Pro Gln Pro Pro Glu Gly Leu Thr Glu
 805 810

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<220>

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23

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23

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29

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cgagccatgc tggctgatgt tc

22

<210> 9
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26

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23

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23

<210> 12

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<211> 25

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25

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23

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ggagctttc ggcaatggct gac

23

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<211> 11

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Tyr Gly Arg Lys Lys Arg Arg Gln Arg Arg Lys
1 5 10

<210> 17

<211> 18

<212> PRT

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<220>

<223> Description of Artificial Sequence: Peptide

<400> 17

Phe Ile Thr Cys Gly Gly Tyr Gly Arg Lys Lys Arg Arg Gln Arg
1 5 10 15

Arg Arg

FIGURE 1

(Map of Human IL-17 Receptor Like cDNA and Amino Acid Sequences)

(SEQ ID NO:1 and SEQ ID NO:2)

ATCATGATCCCTATGGCCTGGCAGTTAGGAATCTCCTGGCTGGCGGGAGTTGACAGCT 60
 61 GTGGGGGGCCATGAAGCGGGCAGAAAGAGAGGTGGATGATGTCCGGGACTGGCATGACC 120
 * S G Q K E R W M M S G D W H D P -
 121 CTGGGTACAGCAGTGCTGCTGCATTGACTCCATGGGCTTGTGGAGAGCAA 180
 G S Q Q C C L H L D S M G L C V G R A N -
 181 ATTGGCTTCACTCTGCATCATGTTCTTTCCCACAGGGAGTGGGCCAGCCAGCA 240
 W L H S A S C S L V F P Q G V G P A S R -
 240 GAAACAGTGGCTGTACAACATCACCTCAAATATGACAATTGTACCACTACTGAATC 300
 N S G L Y N I T F K Y D N C T T Y L N P -
 301 CAGTGGGAAGCATGTGATTGCTGACGCCAGAATATCACCATCAGCCAGTATGCTTGCC 360
 V G K H V I A D A Q N I T I S Q Y A C H -
 361 ATGACCAAGTGGCAGTCACCATTCTTGGCCCCAGGGGCCCTCGGCATCGAATTCTGA 420
 D Q V A V T I L W S P G A L G I E F L K -
 421 AAGGATTCGGTAATACTGGAGGAGCTGAAGTCGGAGGGAAAGACAGTGCCAACA 480
 G F R V I L E E L K S E G R Q C Q Q L I -
 481 TTCTAAAGGATCCGAAGCAGCTAACAGTAGCTTCAAAAGAACTGGAATGGAATCTAAC 540
 L K D P K Q L N S S F K R T G M E S Q P -
 541 CTTCCCTGAATATGAAATTGAAACGGATTATTCGTAAAGGTTGTCCCTTTCTTCCA 600
 F L N M K F E T D Y F V K V V P F P S I -
 601 TTAAAAACGAAAGCAATTACCACCCCTTCTTCTTAAAGGTTGTCCCTTTCTTCCA 660
 K N E S N Y H P F F R T R A C D L L L -
 661 TACAGCCGGACAATCTAGCTTGTAAACCCCTCTGGAGCCTCGGAACCTGAACATCAGCC 720
 Q P D N L A C K P F W K P R N L N I S Q -
 721 AGCATGGCTGGACATGCAGGTGTCTCGACCACGCACCGCACAACTCGGCTTCCGTT 780
 H G S D M Q V S F D H A P H N F G F R F -
 781 TCTTCTATCTCACTACAAGCTCAAGCACGAAGGACCTTCAAGCGAAAGACCTGTAAGC 840
 F Y L H Y K L K H E G P F K R K T C K Q -
 841 AGGAGCAAACACTACAGAGATGACCAAGCTGCTCCTTCAAAATGTTCTCCAGGGATTATA 900
 E Q T T E M T S C L L Q N V S P G D Y I -
 901 TAATTGAGCTGGTGGATGACACTAACACAAGAAAAGTGTGATGCATTATGCCTTAAAGC 960
 I E L V D D T N T T R K V M H Y A L K P -

3721 GGGGGGCTTGCACCCCTGCTGGAAAGAAAAGTTGTATATTCTGATGCAAATGTCA 3780
3781 TACTCACTGCTCTGTAAAGGCAGCTGGCAGCTTTGGAAAAGAACGTGCTCGTCTGTT 3840
3841 CTCTGGCATCAAGTTCTTGCAGCTGCTCTGAGGGAGAGACAGTGAGCTGCAAGACTGCC 3900
3901 TCCCCATAACAACAGGCAACTCAGAGAAGAGTCATTTATGTTGTTCTATGGAATCTGG 3960
3961 AATGAGTGCAGAGCTCCTACCCACACATGACTGCCCGCCATTTCATCCTAGGCATTCTG 4020
4021 TGAAGGAGATTGGTTAGTCCAAACTTGCTAACATACGAAAATTCACTTGGAACATGATGA 4080
4081 GAGATTCTTATTGAGGCCAAGAGATGTTCCCTGTCCCAGAGGAACCATTAGGAGTCGCT 4140
4141 TTAGGGTATTCACTGTTGTTGAAATAAGGCATCTCTGAGAAAGTGGCCCCAGGGAG 4200
4201 AGAATGGAGGACTGGGAGGAGAACATTAAC TGAGCTCCAAGGGTGTGGCAGAGAGC 4260
4261 TTGCTATGTGAACTCACCTTAAGAAAATGGAAGAGAAAAAGAGAGTGCTAGTTAAAAAA 4320
4321 ATCGGGATGTTTAGTTGGATTAGGGTTTGATACTTATGTTGAAATACTAATGTTTC 4380
4381 TGATCAATAAAATCAAACCTTAATATACCGAGTAATGAAACCATA GTGATTGCCTCA 4440
4441 GAATAAATTGAGAAGTCCAAAAAAAAAAAAAAAAAAAAAA 4483

FIGURE 2Overlap of IL-17 Human Receptor-Like (hIL-17RL) Polypeptide and Known Human IL-17 Receptor Family Member

(SEQ ID NO: 3)

hIL-17RL	MMSGDWHDPGSQQCCLHLDMSGLCVGRANLHSASCSDLVFPQGVGPASRN	50
Human IL-17R	..MGAARSPPS...AVPGPLLGLLLLLLGV LAPGGASLRLLDHRALVCSQ	45
hIL-17RL	SGLYNITFKYDNC..TTYLNPVGKHVIADAQNITISQYACHDQ....VA	93
Human IL-17R	PGL. NCTVKNSTCLDDSWIHP. RNLTTPSSPKDLQIQLHFAHTQQGDLFPV	93
hIL-17RL	VTILWS. PGALGIEFLKGFRVILEELKSEGRQCQQLILKDPKQLNSSFKR	142
Human IL-17R	AHIEWTLQTDASILYLEGAELSVLQLNTNERLCVR..FEFLSKLRHHHRR	141
hIL-17RL	TGMESQPFLNMKFETDYFVKV..VPFPSIKNESNYHFFFFRTRACDLLLQ	190
Human IL-17R	WRFTFSHFV.VDPDQEYEVTVHHLPKPIP DGDPNHQSKNFLVPDCEHARM	190
hIL-17RL	PDNLACK...PFWKPR.....NL.NISQHGSDMQVSFDHA	221
Human IL-17R	KVTTPCMSSGSLWDPNITVETLEAHQLRVSTLWNESTHYQILLTSFPHM	240
hIL-17RL	PHNFGFRFFYLHYKLKHEGPFKRKTCKQEQTTEMTSCLLQNVS PGDYII E	271
Human IL-17R	ENHSCFEHMH.HIPAPRPEEFHQRSNVTLRLNKGCCRHQVQIOPFFSS	289
hIL-17RL	LVDDT...NTTRKVMHYALKPVHSPWAGPIRAVAITVPLVVISA FATLFT	318
Human IL-17R	CLNDCLRHSATVSCPEMPDTPEPIP DYMLWVYWFITGISILLVG SVILL	339
hIL-17RL	VMCRKKQQENIYSHLDEESSESSTYTAALPRERLRPRP...KVFLCYSS	364
Human IL-17R	IVCMTWRLAGPGS...EKYSDTDKYTDGLPAADLIPPLKPRKVWIIYSA	386
hIL-17RL	KDGQNHMNVQCFAYFLQDFCGCEVALDLWEDFSLCREGQREWV.IQK..	411
Human IL-17R	.DHPLYDVVLKFAQFLLTAGTEVALDLLEEQAISEAGVMTWVGRQKQE	435
hIL-17RL	.IHESQFIIVVCSKG MK....YFVDKKNYKHKGGGRGSGKGELFLVA VSA	456
Human IL-17R	MVESNSKIIIVLCRGTRAKWQALLGRGAPVRLRCDHGKPVGDLFTAAMNM	485
hIL-17RL	IAEKLQRQAKQSSAALSKFIAVYF.DYSCEGDVPGI LLDLSTKYRIMDNLP	505
Human IL-17R	ILPDFK R....PACFGTYVVCYFSEVSCDGDVPDFGAAPRYPLMDRFE	530
hIL-17RL	QLCSHLHSRDHGLQEPGQHTRQG..SRRNYFRSKSGRSLYVAICNMHQFI	553
Human IL-17R	EV..YFRIQDLEMFPQGRMHRV GELSGDNYLRS PGGRQLRAALDRFRDWQ	578
hIL-17RL	DEEPDWFEKQFV....PFHPPPLR...YREPVL EKDDSLVLNDVMCKPG	596
Human IL-17R	VRCPDW FECENLYSADDQDAPS LDEEV FEEPLLPP.GTGIVKRAPLVRE.	626
hIL-17RL	PESDFCLKVBAAV LGATGPAD SQHE.....SQHGG	626
Human IL-17R	PGSQACLAIDPLVGE EGGA AVAKLEPHLQPRGQPA PQPLH TLV LAEEGA	676

